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Best Local Similarity
Matches 21; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ,29-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of gp41. This sequence which in turn was used 147 (AAB12254).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antigenic peptides and peptide functional derivatives, useful for detection of antibodies produced in response to human immunodeficiency virus group O antibodies -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                De Leys R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-NOV-1999;
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08-FEB-1999;
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                                                                                                                                         Human immunodeficiency virus type 1; HIV-1; envelop; group M; group O; C2V3-env; gp41; gag; retrovirus; strain; gp120; hypervariable loop; primer; hybridisation; amplification; PCR; polymerase chain reaction;
                                                                                                                                                                                                                                                                                                            03-JUN-1997
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  WO9627013
                                                                                                                                                                                                                                                     Partial sequence of gp41 from HIV-1 gp. O strain BCF02 (ESS).
                                                                                                                                                                                                                                                                                                                                                                                                                           AAW07343 standard;
                                                             Human immunodeficiency virus type 1.
                                                                                                                 .mmunogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12
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etliqnqqllnswgckgrivcytsvkwn
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99US-0119138.
99US-0433428.
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                                                                                                                                                                                                                                                                                                                                                                                                                        peptide;
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75.0%;
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Pred. No. 8.2e-12;
5; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                  RESULT
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Best Local S
Matches 21
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N-PSDB; AAT44918.
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                                                                                                                                                                            Group O human immune deficiency virus; HIV; detection; infection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New strains of HIV-1
              24-FEB-1998;
09-APR-1997;
                                                      06-APR-1998;
                                                                                15-OCT-1998.
                                                                                                                                                  Synthetic
                                                                                                                                                                                                     Peptide derived
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                                                                                                                                     deficiency virus
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21; Conservative
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                                                                                                                                                                                                                                 (first entry)
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              98FR-0002212
97FR-0004356
                                                      98WO-FR00691
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75.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 131;
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Best Local
                                                                                                                    The present sequence is peptide 147 of Human Immunodeficiency Virus Type 1 (HIV-1). This sequence is a partial consensus sequence of the immunodominant region of gp41 protein derived from a variety of HIV-1 group O (outlier) strains: ANT70, MVP5180, VAU, DUR, POC, FAN, LOB, MAN, NAN, ESS, NKO, BCF09, BCF12, BCF13, BCF14, 686, ABT063, ABT124, ABT1123, ABT2156, 193Ha, CDC7755, CDC1897, HLD28, 1515, 1516, D47-2d, HCYT2C, Nr42 and PE41 (see AAB12207 to AAB12236). HIV is the principle aetiological agent for acquired immunodeficiency syndrome (AIDS). 9p41 is a HIV envelope protein, and so the present sequence may be used as an antigen for the detection of antibodies produced in response to HIV
                                                                                                                                                                                                                                                                                                                                                                                                               30-NOV-1998;
08-FEB-1999;
04-NOV-1999;
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                                                                                                                                                                                                                                                                                       virus
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detection of antibodies pro
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                                                                                     Sequence
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                        l Similarity
22; Conserv
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immunodeficiency syndrome;
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                        Conservative
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iciency syndrome; group 0 HIV; gp41.
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95.78;
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95.7%;
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                                                                                                                                                                                                                                                                                              peptide functional derivatives, useful for oduced in response to human immunodeficiency
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                       Score 133; DE Pred. No. 2.9e 1; Mismatches
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                                  DB 21;
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08-FEB-1999;
04-NOV-1999;
                                                                                                                                                                                                                                                            The present sequence is a peptide 147 related peptide from Human Immunodeficiency Virus Type 1 (HIV-1). This sequence is a partial consensus sequence of the immunodominant region of 9p41 protein derived from a variety of HIV-1 group O (outlier) strains: ANT70, MVP5180, VAU, DUR, PCC, FAN, LOB, MAN, NAN, ESS, NKO, BCF09, BCF12, BCF13, BCF14, 686, ABT063, ABT124, ABT1256, 1934a, CDC7755, CDC1897, HLD28, 1515, LB16, D47-2d, HCYT2c, NK42 and PE41 (see AAB12207 to AAB12236). HIV is the principle aethological agent for acquired immunodeficiency syndrome
        AAB12236 standard; peptide;
                                                                                                                                                                                                                        (AIDS). gp41 is a HIV envelope protein, and so the present sequenched used as an antigen for the detection of antibodies produced in response to HIV infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
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Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ORTH-) ORTHO-CLINICAL DIAGNOSTICS INC
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                                                                           N
                                                                                                 9 QQRLNSWGCKGRIICYTSARWH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Н
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               qqrlnswgckgriicytsarwh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000-402205/35
                                                                                                                                l Similarity
22; Conser
                                                                                                                                                                                                 23 AA;
                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zheng J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human immunodeficiency virus type 1; antibody detection; unodeficiency syndrome; group O HIV; gp41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98US-0110292.
99US-0119138.
99US-0433428.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99EP-0309491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "Any natural amino acid apart from L-asparagine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide;
                                                                                                                                         76.3%;
           33
                                                                           23
                                                                                                   30
                                                                                                                              0;
                                                                                                                                         Score 132; DB 21;
Pred. No. 4e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23
           ĄΑ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ā
                                                                                                                                Mismatches
                                                                                                                                                                                                                                                    and so the present sequence may
                                                                                                                              0;
                                                                                                                                                       Length 23;
                                                                                                                                Indels
                                                                                                                              0
                                                                                                                             Gaps
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RESULT
AAB12257
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                         30-NOV-1998;
08-FEB-1999;
04-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human Immunodeficiency Virus (HIV) is the principle aetiological agent for acquired immunodeficiency syndrome (AIDS). 9941 is a HIV envelope protein, and may be used as an antigen for the detection of antibodies produced in response to HIV infection. Mosaic 9941 proteins were constructed, in which the immunodominant region of group M HIV was replaced by the corresponding region from group O HIV. The mosaic 9941 proteins would be useful as antigens, used in the detection of anti-group O HIV antibodies produced in response to HIV infection. The present seguence is a dihydrofolate reductase (DHFR) fusion protein of one such
                                                                                                                                                                                                                                                                                                                                       HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antigenic peptides and peptide functional derivatives, useful for detection of antibodies produced in response to human immunodeficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              De Leys
  New antigenic detection of a
                                                                                                                                                                                                                                                                           Human
                                                                                                                                                                                                                                                                                                 acquired
                                                                                                                                                                                                                                                                                                                                                                                                         AAB12257 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mosaic protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 5; Fig 6; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-402205/35
                                     WPI; 2000-402205/35.
                                                            De Leys
                                                                                                                                                        29-NOV-1999;
                                                                                                                                                                                28-JUN-2000
                                                                                                                                                                                                      EP1013766-A2
                                                                                                                                                                                                                                         Misc-difference
                                                                                                                                                                                                                                                                                                               HIV-1;
                                                                                                                                                                                                                                                                                                                                                             10-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                      AAB12257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                   (ORTH-) ORTHO-CLINICAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ORTH-) ORTHO-CLINICAL DIAGNOSTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
les 27; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           w
                                                                                                                                                                                                                                                                                                                                                                                                                                   7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   etlmqnqqrlnswgckgriicytsarwh 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ETLMQDQQRLNSWGCKGRIICYTSARWH 30
                                                                                                                                                                                                                                                                         immunodeficiency virus
                                                                                                                                                                                                                                                                                                             AIDS; human immunodeficiency virus
                                                            ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              æ
                                                                                                                                                                                                                                                                                                 immunodeficiency syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              439 AA;
                                                                                                                                                                                                                                                                                                                                    immunodominant region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         O antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zheng
                                                            Zheng J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
 antibodies produced in
                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                         98US-0110292.
99US-0119138.
99US-0433428.
                                                                                                                                                       99EP-0309491
                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                              /note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90.8%;
                                                                                    DIAGNOSTICS
                                                                                                                                                                                                                           "Any natural amino
                                                                                                                                                                                                                                                                                                                                                                                                           28
                                                                                                                                                                                                                                                                          type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 157; DB 21;
Pred. No. 2.4e-14;
                                                                                                                                                                                                                                                                                                                                                                                                           ₿
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                   consensus sequence peptide 147
  functional derivatives, n response to human immu
                                                                                    INC
                                                                                                                                                                                                                                                                                                 virus type 1; antibody group O HIV; gp41.
                                                                                                                                                                                                                            acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                           apart from L-asparagine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
ves, useful for immunodeficiency
                                                                                                                                                                                                                                                                                                              detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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w
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3.

The present sequence is peptide 147-4 from Human Immunodeficiency Virus Type 1 (HIV-1). This sequence is a partial consensus sequence of the immunodominant region of gp41 protein derived from a variety of HIV-1 group O (outlier) strains: ANT70, MVP5180, VAU, DUR, POC, FAN, LOB, MAN, ESS, NKO, BCF09, BCF12, BCF13, BCF14, 686, AB7063, AB7124, AB71123, AB72156, 193Ha, CDC7755, CDC1897, HLD28, 1515, 1516, D47-2d, HCYT2c, Nr42 and PE41 (see AAB12207 to AAB12236). HIV is the principle aetiological agent for acquired immunodeficiency syndrome (AIDS). gp41 1

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RESULT
AAB1225
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                               30-NOV-1998;
08-FEB-1999;
04-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunodeficiency Virus Type 1 (HIV-1). This sequence is a partial consensus sequence of the immunodominant region of gp41 protein derived from a variety of HIV-1 group 0 (outlier) strains: ANT70, MVP5180, VAU, DUR, POC, FAN, LOB, MAN, NAN, ESS, NKO, BCF09, BCF12, BCF13, BCF14, 686, ABT063, ABT124, ABT1123, ABT2156, 193Ha, CDC7755, CDC1897, HLD28, 1515, 1516, D47-2d, HCYT2c, Nr42 and PE41 (see AAB12207 to AAB12236). HIV is the principle aetiological agent for acquired immunodeficlency syndrome (AIDS), gp41 is a HIV envelope protein, and so the present sequence may be used as an antigen for the detection of antibodies produced in response to HIV infection.
                                                                                                    Claim 1; Page 36; 52pp; English.
                                                                                                                           virus group O antibodies
                                                                                                                                       New antigenic peptides and peptide detection of antibodies produced
                                                                                                                                                                            HPI;
                                                                                                                                                                                                 De Leys
                                                                                                                                                                                                                                                                                                                                              EP1013766-A2
                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency
                                                                                                                                                                                                                                                                                                                                                                                         HIV-1; AIDS; acquired imm
                                                                                                                                                                                                                                                                                                                                                                                                                             HIV-1 gp41 immunodominant region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB12256;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB12256 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is a peptide 147 related peptide from Human Immunodeficiency Virus Type 1 (HIV-1). This sequence is a partial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           virus group O antibodies
                                                                                                                                                                                                                        (ORTH-) ORTHO-CLINICAL DIAGNOSTICS
                                                                                                                                                                                                                                                                                               29-NOV-1999;
                                                                                                                                                                                                                                                                                                                       28-JUN-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ω
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 96.4
27; Conservative
                                                                                                                                                                                                 R, Zheng J;
                                                                                                                                                                                                                                                                                                                                                                                         immunodeficiency
immunodeficiency syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA;
                                                                                                                                                                                                                                             98US-0110292.
99US-0119138.
99US-0433428.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     89.68;
                                                                                                                                                                                                                                                                                                                                                                   virus type
                                                                                                                                                    peptide functional derivatives,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 155; DB 21;
Pred. No. 2.5e-15;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                            consensus sequence peptide
                                                                                                                                                                                                                                                                                                                                                                                                      virus
                                                                                                                                                                                                                                                                                                                                                                                           group
                                                                                                                                        response to
                                                                                                                                                                                                                                                                                                                                                                                            type 1; antibody O HIV; gp41.
                                                                                                                                       erivatives, useful for human immunodeficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28;
                                                                                                                                                                                                                                                                                                                                                                                                       detection;
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                                                                                                                                                                                                                                                                                                                                                                                                                               147-4.
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Fri

Jun

RESULT AAB12261

4

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RESULT
AABIZES3
ID AABI
XX AABI
AC AABI
XZ DIFF
XX HIV-
KW HIV-
KW acqu
                                                                                                                                                                                                                                                                                                               ρ
                                                                                                                                                                                               В
                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-NOV-1998;
08-FEB-1999;
.04-NOV-1999;
                                                                                                                                                                                                                                                                                                                                               Human Immunodeficiency Virus (HIV) is the principle aetiological agent for acquired immunodeficiency syndrome (AIDS). gp41 is a HIV envelope protein, and may be used as an antigen for the detection of antibodies produced in response to HIV infection. Mosaic gp41 proteins were constructed, in which the immunodominant region of group M HIV was replaced by the corresponding region from group O HIV. The mosaic gp41 proteins would be useful as antigens, used in the detection of anti-group O HIV antibodies produced in response to HIV infection. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
             acquired
                          HIV-1; AIDS;
                                                  DHFR-hES-MH fusion protein
                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 15; Page 37-38; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              acquired
                                                                             10-NOV-2000
                                                                                                                               AAB12253 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            detection of antibodies produced in response to human immunodeficiency virus group O antibodies -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antigenic peptides and peptide functional derivatives, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        De Leys R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ORTH-) ORTHO-CLINICAL DIAGNOSTICS INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV group M/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB12261 standard;
                                                                                                        AAB12253;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HIV-1; AIDS;
                                                                                                                                                                                               44
                                                                                                                                                                                                                        ω
                                                                                                                                                         თ
                                                                                                                                                                                             etlmqnqqrlnswgckgriicytsarwh 71
                                                                                                                                                                                                                       ETLMQDQQRLNSWGCKGRIICYTSARWH 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-402205/35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IDS; human immunod immunodeficiency
             immunodeficiency syndrome;
                                                                                                                                                                                                                                                                                                                                       is one such
                                                                                                                                                                                                                                                                                                               220 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zheng J;
                                                                                                                                                                                                                                                Conservative
            human immunodeficiency virus type 1; DHFR-hES-MH; unodeficiency syndrome; group O HIV; gp41.
                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0110292.
99US-0119138.
99US-0433428.
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iciency syndrome; group O HIV; gp41.
                                                                                                                               peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     O mosaic protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide; 220 AA
                                                                                                                                                                                                                                                           90.8%;
                                                                                                                                                                                                                                                                                                                                       mosaic protein.
                                                                                                                                                                                                                                                μ;
                                                                                                                                 368
                                                                                                                                                                                                                                               Score 157; DB
Pred. No. 1.1e
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      type 1.
                                                                                                                                ΑA
                                                                                                                                                                                                                                                           DB 21;
.1e-14;
                                                                                                                                                                                                                                                                       Length 220;
                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                Gaps
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Вb
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Best Local S
Matches 27
30-NOV-1998;
08-FEB-1999;
04-NOV-1999;
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08-FEB-1999;
04-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Immunodeficiency Virus (HIV) is the principle aetiological agent for acquired immunodeficiency syndrome (AIDS). gp41 its a HIV envelope protein, and may be used as an antigen for the detection of antibodies produced in response to HIV infection. Mosaic gp41 proteins were constructed, in which the immunodominant region of group M HIV was replaced by the corresponding region from group O HIV. The mosaic gp41 proteins would be useful as antigens, used in the detection of anti-group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antigenic peptides and peptide detection of antibodies produced virus group O antibodies -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus type 1. Unidentified.
                                                                                                               Human immunodeficiency virus type Unidentified.
                                                                                                                                                acquired
                                                                                                                                                 HIV-1; AIDS; human immunodeficiency virus type 1; DHFR-hENV-MH; acquired immunodeficiency syndrome; group O HIV; gp41.
                                                                                                                                                                                  DHFR-hENV-MH
                                                                                                                                                                                                                                AAB12252;
                                                                                                                                                                                                                                                      AAB12252 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                           EP1013766-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                      mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                             O HIV antibodies produced in response to HIV infection. The present sequence is a dihydrofolate reductase (DHFR) fusion protein of one
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        De Leys R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP1013766-A2
                                              29-NOV-1999;
                                                                    28-JUN-2000.
                                                                                                                                                                                                          10-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ORTH-) ORTHO-CLINICAL DIAGNOSTICS
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                                                                                                                                                                                                                                                                                                             240
                                                                                                                                                                                                                                                                                                          6
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                                                                                                                                                                                                                                                                                                                                                                                                                                      protein.
                                                                                                                                                                                                                                                                                                                                                         Similarity
27; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                 368 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zheng J;
                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                  tusion protein.
                                                                                                                                                                                                          (first
 98US-0110292.
99US-0119138.
99US-0433428.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0110292.
99US-0119138.
99US-0433428.
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                                                                                                                                                                                                                                                      peptide;
                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                   90.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide functional derivatives, oduced in response to human immur
                                                                                                                                                                                                                                                       439
                                                                                                                                                                                                                                                                                                                                                                   Score 157; DB 2
Pred. No. 2e-14;
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                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                DB 21;
                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                              Length 368;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  one such
                                                                                                                                                                                                                                                                                                                                                         Gaps
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Claim 1;

Page

36;

52pp;

English

The present sequence is peptide 147-5 from Human Immunodeficiency Virus Type 1 (HIV-1). This sequence is a partial consensus sequence of the immunodominant region of 9p41 protein derived from a variety of HIV-1 group 0 (outlier) strains; ANTDO, MVP5180, VAU, DUR, POC, FAN, LOB, MAN, ESS, NKO, BCF09, BCF12, BCF13, BCF14, 686, ABT063, ABT124, ABT1123, ABT2156, 193Ha, CDC7755, CDC1897, HLD28, 1515, 1516, D47-2d, HCYT2C, aetLological agent for acquired immunodeficiency syndrome (AIDS). 9p41 is a HIV envelope protein, and so the present sequence may be used as an antigen for the detection of antibodies produced in response to HIV

is

Query Match Best Local

100.0%;

Length

Sequence

30

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RESULT
AAB12259
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The present sequence is a group O Human Immunodeficiency Virus Type 1 (HIV-1) 9p41 peptide. HIV is the principle aetiological agent for acquired immunodeficiency syndrome (AIDS). gp41 is a HIV envelope protein. The present sequence was derived from a sequence homology alignment of group O and group M HIV gp41 descending helix sequences. The present sequence was used as a group O replacement peptide, where was used to construct mosaic gp41 proteins, in which the group M manunodominant region was replaced by the present sequence. The mosaic gp41 proteins (AAB12261 and AAB12262) would be useful as antigens, which
                                                                                                                                                                     New antigenic peptides and peptide functional derivatives, useful for detection of antibodies produced in response to human immunodeficiency virus group O antibodies -
                                                                                                                                         Claim 1; Page 37; 52pp; English.
                                                                                                                                                                                                                                WPI; 2000-402205/35.
                                                                                                                                                                                                                                                          De Leys
                                                                                                                                                                                                                                                                                                                             30-NOV-1998;
08-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                 28-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           {
m HIV-1}; AIDS; human immunodeficiency virus type 1; antibody detection; acquired immunodeficiency syndrome; group O {
m HIV}; {
m gp41}.
                                                                                                                                                                                                                                                                                  (ORTH-) ORTHO-CLINICAL DIAGNOSTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB12259 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                       29-NOV-1999;
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99US-0119138.
99US-0433428.
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Pred. No. 6.9e-18;
); Mismatches 0;
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44 etlmqnqqrlnswgckgriicytsarwh

71

ETLMQDQQRLNSWGCKGRIICYTSARWH 30

Query Match Best Local S Matches 27

27; Conser

Conservative

96.4%;

Score 157; Pred. No. 7

5e-15; DB 21;

Length 149;

Mismatches

Indels

0;

0

Sequence

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RESULT
AAB12262
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Best Local
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               Human Immunodeficiency Virus (HIV) is the principle aetiological agent for acquired immunodeficiency syndrome (AIDS). gp41 is a HIV envelope protein, and may be used as an antigen for the detection of antibodies produced in response to HIV infection. Mosaic gp41 proteins were constructed, in which the immunodominant region of group M HIV was replaced by the corresponding region from group O HIV. The mosaic gp41 proteins would be useful as antigens, used in the detection of anti-group O HIV antibodies produced in response to HIV infection. The present sequence is one such mosaic protein.
                                                                                                                                                                                                                                                                                              30-NOV-1998;
08-FEB-1999;
04-NOV-1999;
                                                                                                                                                Claim
                                                                                                                                                                     New antigenic peptides and peptide functional derivatives, useful for detection of antibodies produced in response to human immunodeficienc virus group O antibodies -
                                                                                                                                                                                                                                              De Leys R,
                                                                                                                                                                                                                                                                                                                                                                                                                       Human
                                                                                                                                                                                                                                                                                                                                                29-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                      28-JUN-2000.
                                                                                                                                                                                                                                                                                                                                                                                               EP1013766-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                               acquired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIV group M/
                                                                                                                                                                                                                                                                       (ORTH-) ORTHO-CLINICAL DIAGNOSTICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIV-1; AIDS;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB12262;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB12262 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        would be used for the detection of anti-group O HIV antibodies produced in response to HIV infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15; Page 38-39; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                     immunodeficiency virus type 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity 96.4
27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                              immunodeficiency
149 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     human immunodeficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 group O mosaic protein
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99US-0119138.
99US-0433428.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            syndrome;
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Pred. No. 1
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                                                                                                                                                                               response to human immunodeficiency
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group O HIV;
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.6e-15;
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Run on: OM protein -

protein search, using sw model

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd

June 20, 2002, 15:26:17; Search time 29.93 Seconds (without alignments)
111.334 Million cell updates/sec

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Minimum DB
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Perfect score:
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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173
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AAB12264
AAB12259
AAB12261
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HIV 9roup M/ 9roup HIV 9roup M/ 9roup DHFR-hES-MH fusion DHFR-hENV-MH fusion HIV-1 9p41 immunod HIV-1 9p41 i
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126	126	127	127	127	127	127	127	127	127	127	127	128	128	128	128	128	128	128	128	128	128	128	128	128	128	128	129	129	130	130	130	130	131	
	72.8				ω ·	73.4	73.4	ω.	ω	ω.		74.0									74.0	•					74.6		٠	•	75.1	75.1	75.7	
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AAW07345	·w	AAY05561	AAW07352	AAB62597	AAY55786	AAY67615	AAY67613	AAY79777	AAY30515	AAB12212	AAG65201	AAW93072	AAR51687	AAY22909	AAW69320	AAB62574	AAY55784	AAY67607	AAY31617	AAW93073	AAR51690	AAB12219	AAB12218	AAB12216	AAB12215	AAB12208	AAW07346	AAB12231	AAY05554	AAY67606	AAB12229	AAW80469	AAW07343	
Partial Sequence o	ial sequence	roup o is	I sequen		- 0	Cide #15 IC	#13 for	ection r	Allelic peptide io	sequence		HIV isolate MVP-51	MVP-5180/91 Iragme	IU NO.	_	9p4 I	gp41 an	`		isolate		Partial sequence o	sequence	sequence	Partial sequence o	Partial sequence o		sequence	group O isc	de #6 for	sequenc	• W		

ALIGNMENTS

RESULT AAB12264

AAB12264 standard; peptide;

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10-NOV-2000 AAB12264;

(first entry)

PTT PTT 30-NOV-1998; 08-FEB-1999; 04-NOV-1999; ${
m HIV}\text{-}1;~{
m AIDS};~{
m human~immunodeficiency~virus~type~1;~antibody~detection;}$ acquired immunodeficiency syndrome; group O HIV; gp41. HIV-1 gp41 immunodominant region consensus sequence peptide 147-5. De Leys R, (ORTH-) ORTHO-CLINICAL DIAGNOSTICS INC 29-NOV-1999; 28-JUN-2000 EP1013766-A2 Human immunodeficiency virus type 1. Zheng J; 98US-0110292. 99US-0119138. 99US-0433428. 99EP-0309491

New antigenic peptides and peptide functional derivatives, useful for detection of antibodies produced in response to human immunodeficiency virus group O antibodies – $\,$

WPI; 2000-402205/35.

2002

WPI; 1998-583190/49.

Delagneau

Gadelle SJX,

Rieunier

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(SNFI) PASTEUR SANOF ,

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RESULT 1
AAB12229
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                                                                                                                                                                                                                     30-NOV-1998;
08-FEB-1999;
04-NOV-1999;
       The present sequence is a partial gp41 protein of Human Immunodeficiency Virus Type 1 (HIV-1) Strain HLD28. HIV is the principle actiological agent for acquired immunodeficiency syndrome (AIDS). gp41 is a HIV envelope protein, and may be used as an antigen for the detection of antibodies produced in response to HIV infection. HLD28 is a member of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AW80459-74 represent synthetic peptides (either linear or cyclised cys-Cys disulphide bonds). The peptides represent variable sequences connected around short highly conserved sequences present in isolate of group O human immune deficiency virus (HIV). The peptides are useful as immunological reagents for detecting infection by group O
                                                                                                         New antigenic peptides and peptide functional derivatives, useful for detection of antibodies produced in response to human immunodeficiency virus group O antibodies -
                                                                                                                                                                                                                                                                                                                   EP1013766-A2
                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus type
                                                                                                                                                                                                                                                                                                                                                               acquired
                                                                                                                                                                                                                                                                                                                                                                          HIV-1; AIDS; human
                                                                                                                                                                                                                                                                                                                                                                                                Partial sequence
                                                                              Example 1;
                                                                                                                                                                         De Leys R,
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                                                                                                                                                                                                 (ORTH-) ORTHO-CLINICAL DIAGNOSTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New synthetic peptide(s) - useful for, human immune deficiency virus of group
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immune deficiency virus (HIV).
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                                                                                                     group O
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                               immunodeficiency syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 44;
                                                                              Fig 1; 52pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                     antibodies
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99US-0119138.
99US-0433428.
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                                                                                                                                                                                                                                                                                                                                                                                                  of HIV-1
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                                                                                                                                                                                                                                                                                                                                                              immunodeficiency virus type 1; immunodominant region;
iciency syndrome; group O HIV; gp41; HLD28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide;
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71.48;
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                                                                                                                                                                                                                                                                                                                                                                                                strain HLD28 gp41 immunodominant region
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Pred. No. 1
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group 0

(outlier).

The present sequence is the

immunodominant region

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     CC infection. The invention relates to peptides derived from HIV-1 which have been modified for use as diagnostic antigens in the treatment or prevention of infection. The structural stability of the peptides can be increased in four different ways; through the replacement of a Nydrophobic amino acid with a less hydrophobic amino acid; through the removal of a positive charge from the peptide, or increase in the amount of secondary structure (i.e. alpha helix) in the constraint of the epitopic sequence via the formation of a covalent crosslink. Modified peptides of the invention are used to detect infectious agents specifically HIV-1. Other detectable agents include Group O viruses; human T-cell lymphotropic virus-I or -II; hepatitis C and the causative agent of syphilis. The peptides can be used for prevention or treatment of infections (e.g. as vaccines, or where cexpressed from a transgene). More generally almost any peptide can be similarly modified, e.g. cytokines and interferons; major this complex antigens; hormones; growth factors; tumour considered complex antigens from many other pathogens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT
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Best Local S
Matches 21
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01-SEP-1998;
15-SEP-1998;
28-JAN-1999;
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which in
                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                            Claim 18;
                                                                                                                                                                                                                                                                                                                     Improving properties of peptides for use preventing or treating infections -
                                                                                                                                                                                                                                                                                                                                                                                           Chowdhury
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptide #6 for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY67606 standard;
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                                                                                                                                                                                                                                                envelope protein,
                                                                                                                                                                                                                                                             peptide is derived from the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    etlmqnqqrldlwgckgrlicytsvkwn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence was used in a sequence homology alignment, turn was used to derive a consensus sequence peptide: pe
                                                                                                                                                                                                                                                                                                                                                                                          MA,
                                                                                                                                                                                                                                                                                          Page 61; 83pp;
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98US-0100422.
99WO-US01726.
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Matches 21; Conservative 5; Mismatches 2; Indels 0; Gaps 0;

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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/2/1aa/5A_COMB.pep:*
2: /cgn2_6/ptodata/2/1aa/5B_COMB.pep:*
3: /cgn2_6/ptodata/2/1aa/6A_COMB.pep:*
4: /cgn2_6/ptodata/2/1aa/6B_COMB.pep:*
5: /cgn2_6/ptodata/2/1aa/FCTUS.COMB.pep:*
5: /cgn2_6/ptodata/2/1aa/Dackfiles1.pep:*
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Gapop 10.0 , Gapext 0.5
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US-09-433-428D-12
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						Sequence 38, Appl			. 68			Sequence 46, Appl				Sequence 10, Appl	Sequence 10, Appl

ALIGNMENTS

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SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 69

LENGTH: 30

TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description
OTHER INFORMATION: Sequence
US-09-433-428D-69
                                     GENERAL INFORMATION:
APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF
FILE REFERENCE: CDS-207
CURRENT APPLICATION NUMBER: US/09/433,428D
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 63
LENGTH: 30
                                                                                                                                                                                                                                       US-09-433-428D-63; Sequence 63, Application US/09433428D; Patent No. 6149910
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/433,428D CURRENT FILING DATE: 1999-11-04 NUMBER OF SEQ ID NOS: 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: De Leys, Rol
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEI
FILE REFERENCE: CDS-20
LENGTH: 30
TYPE: PRT'
ORGANISM: Artificial Sequence
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Pred. No. 2.2e-19;
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; OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus; OTHER INFORMATION: Sequence US-09-433-428D-66
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OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus
OTHER INFORMATION: Sequence
US-09-433-428D-67
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                                                                                                                                                             FILE REFERENCE: CDS-207
CURRENT APPLICATION NUMBER: US/09/433,428D
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 66
LENGTH: 220
                                                                                                                                                                                                                                                                                                                          Sequence 66, Application US/09433428D Patent No. 6149910 GENERAL INFORMATION:
Query Match
Best Local :
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Best Local Similarity
Matches 27; Conserv
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LENGTH: 149
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/433,428D
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES
FILE REFERENCE: CDS-207
                                                                                                                                                                                                                                                                                          APPLICANT: De Leys, Robert J
APPLICANT: Zheng, Jian
                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                 FEATURE:
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Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                         44 ETLMQNQQRLNSWGCKGRIICYTSARWH 71
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29; Conserv
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Pred. No. 2.4e-18;
0; Mismatches 1
 Score
Pred.
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Pred. No. 3.1e-16
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 157;
No. 4.
 DB 4;
.8e-16;
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               Length 220
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; OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus; OTHER INFORMATION: Sequence US-09-433-428D-58
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US-09-433-428D-58
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APPLICANT: De Leys, Rot
APPLICANT: Zheng, Jian
                                                         Query Match
Best Local Similarity
Matches 27; Conserv
                                                                                                                                                                                                                                          SEQ ID NO 57
                                                                                                                                                                                                                                                                                                                                                                                         Sequence 57, Application US/09433428D Patent No. 6149910
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APPLICANT: Zheng, Jian
                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn Ve
                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
FILE REFERENCE: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
FILE REFERENCE: CD:
                                                                                                                                                                                         LENGTH: 439
TYPE: PRT
ORGANISM: Artificial Sequence
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TYPE: PRT
TYPE: PRT
TYPE: Artificial Sequence
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240 ETLMQNQQRLNSWGCKGRIICYTSARWH
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                  3 ETLMQDQQRLNSWGCKGRIICYTSARWH 30
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                                                                                                                                                                                                                                                                                                                                                            De Leys, Robert J
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                                                           Conservative
                                                                                                                                                                                                                                                                                                               CON: PEPTIDES FOR THE DETECTION CDS-207
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                                                                                                                                                                Description of Artificial Sequence: HIV Consensus
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                                                                                                                                                                                                                                                                      1999-11-04
: 70
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                                                                       90.8%;
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                                                         Score 157; DB 4
Pred. No. 1e-15;
1; Mismatches
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 267
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                                                                                     Length 439;
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US-09-433-428D-62

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APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTI
TITLE OF INVENTION: PEPTIDES FOR THE DETECTI
TITLE OF INVENTION NUMBER: US/09/433,428D
CURRENT FILING DATE: 199-11-04
RUMBER OF SEO ID NOS: 70
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 62
LENGTH: 28
LENGTH: 28
TYPE: PRT
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OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus; OTHER INFORMATION: Sequence
US-09-433-428D-62
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US-09-433-428D-64
; Sequence 64, Application US/09433428D
; Patent No. 6149910
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                                                                                                                                                                                                                                                                                                       NAME/KEY: Xaa is any amino acid
LOCATION: 13
OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus
OTHER INFORMATION: Sequence
US-09-433-428D-64
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LENGTH: 35
TYPE: PRT
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US-09-433-428D-61
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Sequence 61, Application US/09433428D Patent No. 6149910 GENERAL INFORMATION: APPLICANT: De Leys, Robert J. APPLICANT: Zheng, Jian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ETLE REFERENCE: CDS-207
CURRENT APPLICATION NUMBER: US/09/433,428D
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP
FILE REFERENCE: CDS-207
FILE REFERENCE: CDS-207
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Best Local Similarity
Matches 27; Conserv
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pred. No. 1e-16;
                                                                                                                                                                                                                                                Score 155; DB 4;
pred. No. 1.3e-16;
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OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus; OTHER INFORMATION: Sequence US-09-433-428D-61
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CURRENT APPLICATION NUMBER: US/09/433,428D
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 70
SOCTWARE: PATENTIN VET: 2.0
SEQ ID NO 61
LENGTH: 23
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US-09-433-428D-59
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OTHER INFORMATION:
OTHER INFORMATION:
CONTROL OF THE CONT
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CURRENT APPLICATION NUMBER: US/09/433,428D
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 59
LENGTH: 23
TYPE: PRT
ORGANISM: Artificial Sequence
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US-09-433-428D-60
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Best Local Similarity
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APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
APPLICANT: Zheng, Jian
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Best Local Similarity
Matches 22; Conserv
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                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: De Leys, Roba
APPLICANT: Zheng, Jian
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEP
FILE REFERENCE: CDS-207
SOFTWARE: P.
SEQ ID NO 60
LENGTH: 23
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 60, Application US/09433428D Patent No. 6149910
                                                                                                                                   FILE REFERENCE: CDS-207
CURRENT APPLICATION NUMBER: US/09/433,428D
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEO ID NOS: 70
NUMBER OF SEO ID NOS: 70
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milarity 95.7%;
Conservative
                                                                                                             PatentIn Ver. 2.0
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milarity 95.7%;
Conservative
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                                                                                                                                                                                                                                                                                          PEPTIDES FOR THE DETECTION OF HIV-1 GROUP
                                                                                                                                                                                                                                                                                                                                                             Robert J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Score 133; DB
; Pred. NO. 1.6e
1; Mismatches
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pred. No. 1.1e-13
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.6e-13
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US-08-894-699-36
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 30
LENGTH: 33
TYPE: PRT
ORGANISM: Human immunodeficiency virus type 1
US-09-433-428D-30
                                                                                                                                                                                                                                                                                                                                        Sequence 36 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-433-428D-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus; OTHER INFORMATION: Sequence
US-09-433-428D-60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 30, Application US/09433428D Patent No. 6149910
           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/433,428D CURRENT FILING DATE: 1999-11-04 NUMBER OF SEQ ID NOS: 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES
                                                                                                                                                                                                      APPLICANT: CHAIX-BAUDIER, MARIE-LAURE
TITLE OF INVENTION: GROUP O HIV-1, FRAGMENTS
TITLE OF INVENTION: VIRUSES, AND USES THEREOF
                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                    NUMBER OF SEQUENCES:
   OPERATING SYSTEM:
                                                                                                                     STREET: 1755 STREET: FLOOR
                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: Xaa is any amino acid LOCATION: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                       ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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5. 6030769
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                                                                                                         ARLINGTON
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                                                                                                                         1755 SOUTH JEFFERSON DAVIS HIGHWAY, FOURTH
                                                                              USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         76.3%; Score 132; ilarity 100.0%; Pred. No. Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ON: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP O CDS-207
compatible
PC-DOS/MS-DOS
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US-09-444-410-36
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Best Local Similarity
Matches 21; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 40 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
SOFTWARE: Patentin Release #
                PRIOR APPLICATION DATA:
                                                                                                                     SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: OBLON, NORMAN F.
REGISTRATION NUMBER: 24,614
REFERENCE/DOCKET NUMBER: 03:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
                                    FILING DATE
                                                                                                APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: PCT/I
FILING DATE: 26-FEB-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 95
FILING DATE: 27-FEB-1995
                                                                                                                                                                                                                                      COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11 ETLIQNQQLLNSWGCKGRIVCYTSVKWN 38
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CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 ETLMQDQQRLNSWGCKGRIICYTSARWH 30
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TELEPHONE: 703-413-2220
TD TD NO:
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75.0%;
FR 95/02236
                                           08/894,699
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Pred. No. 5.7e-13;
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                                                                                                                                    Version #1.30
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Sequence 23, Application US/09433428D

Patent No. 6149910

GENERAL INFORMATION:
APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP O
FILE REFERENCE: CDS-207

CURRENT APPLICATION NUMBER: US/09/433,428D

CURRENT FILING DATE: 1999-11-04

NUMBER OF SEQ ID NOS: 70

SOFTWARE: Patentin Ver. 2.0

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 23
LENGTH: 33
TYDE: PRT
ORGANISM: Human immunodeficiency virus type 1

US-09-433-428D-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: OBLOW, NORMAN F.

REGISTRATION NUMBER: 24,614
REFERENCE/DOCKET NUMBER: 0354-0020-0PCT
RELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
TELEPHAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
LENGTH: 40 amino acids
TYPE: amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-09-444-410-36
Scarch completed: June 20, 2002, 15:31:50 Job time: 218 sec
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Best Local Similarity
Matches 21; Conserv
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                        Score 130; DB 4; Length 33; Pred. No. 6.5e-13; 5; Mismatches 2; Indels
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Maximum DB seq length: 2000000000
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                                                                                                                                         Score
   173
173
166
166
157
157
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100.0
96.0
96.0
90.8
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Listing first 45 summaries
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173
                                                                                                                                                                                                                                                                                       : /cgn2_6/ptodatta/2/paa/US081_COMB.pep:*
    /cgn2_6/ptodatta/2/paa/US081_COMB.pep:*
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Copyright (c) 1993 - 2000 Comp
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US-09-605-573A-69
US-09-433-428-63
US-09-605-573A-63
US-09-605-573A-63
US-09-433-428-67
US-09-605-573A-67
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-09-886-150-6	-09-886-149-6	-09-147-362A-	-09-147-362-21	-09-605-573A-1	-09-605-573A-	-09-605-573A-1	-09-605-573A-	-09-605-573A-	-09-433-42	-09-433-428-1	-09-433-428-1	-09-433-428-	-09-433-428-	-09-605-573A-	-09-433-428-2	-09-462-917A-	-09-605-573A-2	-09-433-428-	-09-147-362A-	9-14	-09-605-573A-	-09-433-428-3	-09-605-573A-	-09-433-428-60	-09-605-573A	-09-433-428-5	-09-605-573A-	-09-433-428-6	-09-605-573A	-09-433-428-6	-09-605-573A	-09-433-428-6	-09-605-573A-	-09-433-428-5	S-09-605-573A-	US-09-433-428-58	-09-605-573A-
equence	equence	equence	equence	equence	equence	equence	equence	ø	equence	equence	equence 10	equence	equence	equence	equence	equence	equence	ø	equence	Sequence 11	equence	equence	equence	equence	equence	equence	equence	equence	equence	Ð	equence	equence	equence	equence	equence	quence	equence
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ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/09/433,428
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 69
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 69
LENGTH: 30
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence
US-09-433-428-69
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                                                                                                                                                                                                                                                                                                                                                               Sequence 69, Application US/09433428
GENERAL INFORMATION:
APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP OF FILE REFERENCE: CDS-207
                                                                   Query Match
Best Local S
Matches 30
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                1 GRETLMQDQQRLNSWGCKGRIICYTSARWH 30
GRETLMQDQQRLNSWGCKGRIICYTSARWH
                                                                 th 100.0%; Score 173; DB 1 Similarity 100.0%; Pred. No. 7e-17; 30; Conservative 0; Mismatches
                                                                                                    DB 18;
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                                                                                                    Length 30;
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US-09-605-573A-69
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GENERAL INFORMATION:
APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/433,428
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 69
SOFTWARE: Patentin Ver. 2.0
SEG ID NO 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 63, Application US/09433428 GENERAL INFORMATION:
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CURRENT FILING DATE: 2000-06-28
NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 69
LENGTH: 30
                                                                    Sequence 63, Application US/09605573A
GENERAL INFORMATION:
APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
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Best Local :
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP
FILE REFERENCE: CDS-207
CURRENT APPLICATION NUMBER: US/09/605,573A CURRENT FILING DATE: 2000-06-28
                                   TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF FILE REFERENCE: CDS-222
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                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE: OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: Sequence
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96.7%;
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Pred. No. 6.9e-16;
0; Mismatches 1
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                                                        HIV-1 GROUP
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; FEATURE:
; NAME/KEY: Xaa is any amino acid
; LOCATION: 8
; OTHER INFORMATION: Description of
; OTHER INFORMATION: Sequence
US-09-605-573A-63
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                                               SEQ ID NO 67
LENGTH: 149
TYPE: PRT
ORGANISM: Artificial Sequence
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LENGTH: 30
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Best Local :
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP
FILE REFERENCE: CDS-207
CURRENT APPLICATION NUMBER: US/09/433,428
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 69
NUMBER OF SEQ ID NOS: 69
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                CURRENT APPLICATION NUMBER: US/09/605,573A CURRENT FILING DATE: 2000-06-28 NUMBER OF SEQ ID NOS: 70
                                                                                                                                                                                                                       APPLICANT: De Leys, Robert J. APPLICANT: Zheng, Jian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS:
SOFTWARE: Patentin Ve
                                                                                                                    SOFTWARE:
                                                                                                                                                                                       TITLE OF INVENTION: PEP FILE REFERENCE: CDS-222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 149
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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FEATURE: OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                     Jocal Similarity 96.4%; nes 27; Conservative
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                                                                                                                    PatentIn Ver. 2.0
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96.7%;
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    Mismatches

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Pred. No. 6.9e-16;
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US-09-433-428-66
; Sequence 66, Application US/09433428
; GENERAL INFORMATION:
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CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 69
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 66
LENGTH: 220
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Best Local Similarity 96.4

Matches 27; Conservative
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CURRENT FILING DATE: 2000-06-28
CURRENT FILING DATE: 2000-06-28
COURTENT FILING DATE: 2000-06-28
COURTENT FILING DATE: 200 FALENCTH: 220
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                       Sequence 66, Application US/09605573A GENERAL INFORMATION:
APPLICANT: De Leys, Robert J.
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                                                                   Query Match
Best Local
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TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP O
FILE REFERENCE: CDS-207
                                                                                                                                                                                                                                                                                               FILE REFERENCE: CDS-222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus OTHER INFORMATION: Sequence
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                                                                                                                                                                               ORGANISM: Artificial Sequence
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                                                       Local Similarity
nes 27; Conser
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Conservative
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                                                                   96.4%;
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                                                                   Score 157; DB 20
Pred. No. 9e-14;
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.2e-14;
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US-09-605-573A-58
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US-09-433-428-57
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GENERAL INFORMATION:
APPLICANT: De Leys, Robert J.
APPLICANT: Lheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP O
FILE REFERENCE: CDS-207
                                                                                                  Sequence 57, Application US/09433428 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 27; Conserv
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SEQ ID NO 58
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SEQ ID NO 58
LENGTH: 368
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 58, Application US/09605573A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: PEPTIDES FOR THE DETECTIVE REFERENCE: CDS-222
CURRENT APPLICATION NUMBER: US/09/605,573A
CURRENT FILING DATE: 2000-06-28
NUMBER OF SEQ ID NOS: 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/433,428
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 69
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION OF HIV-1 GROUP OF FILE REFERENCE: CDS-207 CURRENT APPLICATION NUMBER: US/09/433,428 CURRENT FILING DATE: 1999-11-04
                                                                APPLICANT: De Leys, Robert J. APPLICANT: Zheng, Jian
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APPLICANT: Zheng, Jian
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OTHER INFORMATION: Descripti
                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 368
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                                                                                                                                                                                                         240
                                                                                                                                                                                                                                                                          Local Similarity
mes 27; Conserv
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Pred. No. 1.5e
1; Mismatches
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Pred. No. 1.5e-13;
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1.5e-13;
hes 0;
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; OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus; OTHER INFORMATION: Sequence US-09-433-428-62
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Sequence 62, Application US/09433428
GENERAL INFORMATION:
APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: EMPTIDES FOR THE DETECTION OF HIV-1 GROUP O
FILE REFERENCE: CDS-207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus ; OTHER INFORMATION: Sequence US-09-605-573a-57
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Best Local S
Matches 27
                                                                                                  CURRENT APPLICATION NUMBER: US/09/433,428
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 69
SOFTWARE: PATENTIN VEr. 2.0
SEQ ID NO 62
LENGTH: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/605,573A CURRENT FILING DATE: 2000-06-28 NUMBER OF SEQ ID NOS: 70 SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 57 LENGTH: 439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 57, Application US/09605573A GENERAL INFORMATION:
APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR TH
FILE REFERENCE: CDS-222
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Best Local Similarity
Matches 27; Conserv
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SOFTWARE: PatentIn Ver. .
SEQ ID NO 57
LENGTH: 439
                                                                TYPE: PRT
ORGANISM: Artificial Sequence
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                                                FEATURE:
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nes 27; Conserv
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Pred. No. 1.8e
1; Mismatches
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Pred. No. 1.8e
1; Mismatches
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.8e-13;
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.8e-13;
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                                                                                                                                                    US-09-433-428-64
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LENGTH: 35
TYPE: PRT
                                                                Query Match
Best Local S
Matches 27
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Best Local S
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APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES
                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/433,428
CURRENT FILING DATE: 1999-11-04
NUMBER OF SEQ ID NOS: 69
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                     APPLICANT: De Leys, Robert J.
APPLICANT: Zheng, Jian
TITLE OF INVENTION: PEPTIDES FOR THE DETECTION
FILE REFERENCE: CDS-207
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CURRENT APPLICATION NUMBER: US/09/605,573A
CURRENT FILING DATE: 2000-06-28
NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence:HIV Consensus OTHER INFORMATION: Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 28
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                    ORGANISM: Artificial Sequence FEATURE:
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               3 ETLMQDQQRLNSWGCKGRIICYTSARWH 30
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ETLMQXQQRLNSWGCKGRIICYTSARWH
                                                                Similarity 96.4
27; Conservative
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96.4%;
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Pred. No. 3e-1
0; Mismatches
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Pred. No. 2.4e-14;
0; Mismatches 1
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Pred. No. 2.4e-14;
0; Mismatches 1
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                                                                                                 DB 18;
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                                                                                                 Length 35;
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1: /ggn2_6/ptodata/1/paa/PCT_NEW_COMB.pep:*

2: /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:*

3: /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*

4: /ggn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*

5: /cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*

6: /cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*

7: /cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*
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177.533 Million cell updates/sec
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US-09-914-205-2
US-09-913-159x-2
US-09-913-159x-2
US-09-913-159x-2
US-09-915-5-8
US-08-776-188B-19
US-08-776-188B-19
US-08-776-188B-10
US-08-776-188B-54
US-08-776-188B-54
US-08-776-188B-54
US-08-776-188B-54
US-08-776-188B-59
US-08-776-188B-39
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   Sequence 2, Appli
Sequence 23, Appli
Sequence 24, Appli
Sequence 36, Appli
Sequence 8, Appli
Sequence 19, Appli
Sequence 10, Appli
Sequence 20, Appli
Sequence 11, Appli
Sequence 14, Appli
Sequence 5, Appli
Sequence 6, Appli
Sequence 30, Appli
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Sequence 5, Appli
Sequence 3, Appli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1
US-09-914-205-2
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APPLICANT: Rovinski, Benjamin
APPLICANT: ROVINSKI, Benjamin
TITLE OF INVENTION: EXPRESSING GP140 FRAGMENT OF PRIMARY HIV-1 ISOLATE
FILE REFERENCE: 1038-1176 MIS:jb
CURRENT APPLICATION NUMBER: US/09/914,205
CURRENT FILING DATE: 2002-04-09
CURRENT FILING DATE: 2002-04-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 09/256,194
PRIOR FILING DATE: 1999-02-24
NUMBER OF SEQ ID NOS: 17
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51 29.5 275 7 US-60-360-039-4600 Sequence 4500, Ap 51 29.5 275 7 US-60-360-039-7357 Sequence 7357, App 49 28.3 13 5 US-09-576-824A-170 Sequence 110, App 49 28.3 14 5 US-09-576-824A-176 Sequence 177, App 49 28.3 14 5 US-09-576-824A-176 Sequence 176, App 49 28.3 15 5 US-09-576-824A-111 Sequence 9, Appl 14 28.3 15 5 US-09-576-824A-111 Sequence 9, Appl 15 US-09-573-655B-1352 Sequence 111, App 49 28.3 477 5 US-09-573-655B-1352 Sequence 1352, App 16 27, 7 592 7 US-60-360-039-2048 Sequence 23048, Appl 17 200 6 US-10-138-098-38 Sequence 23048, Appl 18 27, 7 1006 6 US-10-138-098-48 Sequence 2688, Appl 18 27, 7 1006 6 US-10-138-098-45 Sequence 45, Appl 18 27, 7 1006 6 US-10-138-098-45 Sequence 47, Appl 18 27, 7 1006 6 US-10-138-098-45 Sequence 47, Appl 18 27, 7 1066 US-10-138-098-45 Sequence 47, Appl 18 27, 7 1066 US-10-138-098-45 Sequence 47, Appl 18 27, 7 1064 Sus-10-138-098-41 Sequence 48, Appl 18 27, 7 1064 Sus-10-138-098-41 Sequence 47, Appl 18 27, 7 1064 Sus-10-138-098-42 Sequence 47, Appl 18 27, 7 1064 Sus-10-138-098-42 Sequence 47, Appl 18 27, 7 1064 Sus-10-138-098-42 Sequence 47, Appl 18 28, Appl 18 28, Appl 18 28, Appl 18 28, Appl 28, Appl 28, Appl 28, Appl 28, Appl 38, Ap	45	44	43	42	41	40	ა 9	3 8	37	36	35 5	34	ω W	32	<u>3</u> 1	30	29	28	27
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7 US-60-360-039-4600 Sequence 4600, 7 US-60-560-24A-110 Sequence 7357, 5 US-09-576-824A-110 Sequence 110, 5 US-09-576-824A-117 Sequence 177, 5 US-09-576-824A-117 Sequence 176, 5 US-09-576-824A-111 Sequence 176, 5 US-09-576-824A-111 Sequence 111, 5 US-09-576-824A-111 Sequence 111, 6 US-09-573-655B-1352 Sequence 135, 7 US-60-360-039-22048 Sequence 2204, 6 US-10-138-098-38 Sequence 2688, 6 US-10-138-098-45 Sequence 45, 6 US-10-138-098-45 Sequence 45, 6 US-10-138-098-45 Sequence 46, 6 US-10-138-098-45 Sequence 47, 6 US-10-138-098-45 Sequence 48, 6 US-10-138-098-45 Sequence 48, 6 US-10-138-098-45 Sequence 48, 7 US-10-138-098-48 Sequence 48, 8 US-10-138-098-49 Sequence 48, 8 US-10-138-098-49 Sequence 48, 8 US-10-138-098-40 Sequence 48, 8 US-10-138-098-41 Sequence 41, 8 US-10-138-098-42 Sequence 42, 8 US-10-138-098-42 Sequence 41, 8 US-10-138-098-42 Sequence 41, 8 US-10-138-098-42 Sequence 42, 8 US-10-138-098-42 Sequence 41, 8 US-10-138-098-42 Sequence 42, 8 US-10-138-098-42 Sequence 41, 8 US-10-138-098-42 Sequence 41, 8 US-10-138-098-43 Sequence 41, 8 US-10-138-098-43 Sequence 41, 8 US-10-138-098-44 Sequence 41, 8 US-10-138-098-44 Sequence 41, 8 US-10-138-098-45 Sequence 41, 8 US-10-138-098-45 Sequence 41, 9 US-10-138-098-14 Sequence 41	27.7	27.7	27.7	27.7	27.7	27.7	27.7	27.7	27.7	27.7	27.7	28.3	28.3	28.3	. 28.3	28.3	28.3	29.5	29.5
US-60-360-039-4600 Sequence 4600 US-60-360-039-7357 Sequence 7357, US-09-576-824A-110 Sequence 110, US-09-576-824A-176 Sequence 177, US-09-576-824A-176 Sequence 176, US-09-576-824A-171 Sequence 176, US-09-571-432-9 Sequence 176, US-09-573-655B-1352 Sequence 1352 US-10-138-098-34 Sequence 34, US-60-360-039-22048 Sequence 2204 US-10-138-098-39 Sequence 39, US-10-138-098-45 Sequence 45, US-10-138-098-45 Sequence 45, US-10-138-098-45 Sequence 45, US-10-138-098-45 Sequence 47, US-10-138-098-46 Sequence 48, US-10-138-098-47 Sequence 47, US-10-138-098-48 Sequence 47, US-10-138-098-41 Sequence 41, US-10-138-098-42 Sequence 41,	1064	1064	1006	1006	1006	1006	849	641	641	592	220	477	15	14	14	13	12	275	275
-60-360-039-4600 Sequence 4500 -60-360-039-7357 Sequence 7357 -09-576-824A-117 Sequence 177, -09-576-824A-176 Sequence 177, -09-576-824A-176 Sequence 177, -09-576-824A-176 Sequence 176, -09-573-655B-1352 Sequence 131, -09-573-655B-1352 Sequence 34, -10-138-098-34 Sequence 34, -10-138-098-38 Sequence 39, -10-138-098-45 Sequence 2688 -10-138-098-45 Sequence 45, -10-138-098-46 Sequence 45, -10-138-098-46 Sequence 45, -10-138-098-47 Sequence 47, -10-138-098-48 Sequence 48, -10-138-098-49 Sequence 47, -10-138-098-49 Sequence 47, -10-138-098-40 Sequence 47, -10-138-098-41 Sequence 47, -10-138-098-42 Sequence 47, -10-138-098-42 Sequence 48, -10-138-098-42 Sequence 47, -10-138-098-42 Sequence 42,	6	0	6	6	0	6	7	6	6	7	6	ر ح	տ	տ	v	ر.	ű	7	7
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ALIGNMENTS

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SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 2

LENGTH: 657

TYPE: PRT

ORGANISM: Human immunodeficiency virus type
US-09-914-205-2
                                                                                                                                                                                                                                                                                                                             US-09-899-575-23
                                                                                                                                                                                                                                                                              Sequence 23, Application US/09899575 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 13; Conserv
                APPLICANT: 2ur Megede, Jan
APPLICANT: Barnett, Susan W.
APPLICANT: Egnelbrecht, Susan W.
APPLICANT: Egnelbrecht, Susan
APPLICANT: van Rensburg, Estrelita Janse
APPLICANT: van Rensburg, Estrelita Janse
TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING ANTIGENIC HIV TYPE C
TITLE REFERENCE: PP01631.102
CURRENT APPLICATION NUMBER: US/09/899,575
CURRENT APPLICATION NUMBER: 09/475,704
PRIOR APPLICATION NUMBER: 09/475,704
PRIOR APPLICATION DATE: 1999-12-30
NUMBER OF SEQ. ID NOS: 135
SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                              565 ERYLODORFLGMWGCSGKLICTTAVPWN 592
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Pred. No. 0.
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0.00033;
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; ORGANISM: Human immunodeficiency virus US-09-899-575-24
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; ORGANISM: Human immunodeficiency virus
US-09-899-575-23
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Query Match
Best Local Similarity
                                                                                                                                        SOFTWARE: P
SEQ ID NO 24
                                                                                                                                                                                                                                                                                                                                                                               Sequence 24, Application US/09899575 GENERAL INFORMATION:
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Best Local Similarity 46.4
Matches 13; Conservative
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LENGTH: 854
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                                                                                                                                                                       CURRENT FILING DATE: 2001-07-05
PRIOR APPLICATION NUMBER: 09/475,704
PRIOR FILING DATE: 1999-12-30
NUMBER OF SEQ ID NOS: 135
                                                                                                                                                                                                                        APPLICANT: van Rensburg, Estrelita Janse
TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING ANTIGENIC
TITLE OF INVENTION: POLYPEPTIDES, POLYPEPTIDES AND USE
FILE REFERENCE: POD1631.102
CURRENT APPLICATION NUMBER: US/09/899,575
CURRENT FILING DATE: 2001-07-05
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LENGTH: 849
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Virus-Vaccine FILE REFERENCE: P057760 CURRENT APPLICATION NUMBER: US/09/913,159A CURRENT FILING DATE: 2002-03-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 199 07 485.2 PRIOR FILING DATE: 1999-02-12 NUMBER OF SEQ ID NOS: 12
                                                                                                      TYPE: PRT
                                                                                                                      LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              582 ERYLKDQQLLGIWGCSGKLICTTAVPWN 609
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13; Conserv
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49.18;
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Score 85;
Pred. No.
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Pred. No. 0.00043;
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DB 5;
0.00043;
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; OTHER INFORMATION: Description of ; OTHER INFORMATION: HIV subtype O US-08-776-188B-8
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                                                                                           SEQ ID NO 8
LENGTH: 15
TYPE: PRT
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APPLICANT:
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Best Local
                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: PCT/EP95/02919
PRIOR FILING DATE: 1995-07-24
PRIOR APPLICATION NUMBER: P 44-30 972.4
PRIOR FILING DATE: 1994-08-31
PRIOR APPLICATION NUMBER: P 44-26 276.0
                                                                                                                                                 PRIOR FILING DATE: 1994-07-25
NUMBER OF SEQ ID NOS: 76
SOFTWARE: PatentIn version 3.
                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/08/776,188B CURRENT FILING DATE: 2002-04-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Weinhues, Ursula-Henrike
APPLICANT: Kruse-Muller, Cornelia
APPLICANT: Hoss, Eva
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                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING MULTIPLE ANTIFILE REFERENCE: 100564-07003
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/130,157
CURRENT FILING DATE: 2002-05-14
PRIOR APPLICATION NUMBER: DE 19955089.1
PRIOR FILING DATE: 1999-11-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: GENEART GmbH
TITLE OF INVENTION: The
FILE REFERENCE: WAG-001
                                                     FEATURE:
                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 379
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/08776188B
                                                                                                                                                                                                                                                                                                                                                                             Wiedmann, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                 Ofenloch-Hahnle, Beatus
                                                                                                                                                                                                                                                                                                                                                                                                                                         Faatz, Elke
                                                                                                                                                                                                                                                                                                                                                                                                   Seidel, Christoph
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Pred. No. 0.
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                           Artificial Sequence: Epitope region of HIV I
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Query Match
Best Local Similarity
Matches 10; Conserv

Conservative

4; Mismatches Score 72; DB 4; Pred. No. 0.00052;

41.6%;

Length 15;

Indels

0;

Gaps

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В
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            12
LSLWGCKGKLVCYTS 15
       LNSWGCKGRIICYTS
             26
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US-08-776-1888-19 GENERAL INFORMATION:
APPLICANT: Weinhues, Ursula-Henrike
APPLICANT: Kruse-Muller, Cornella
APPLICANT: Hoss, Eva
APPLICANT: Foatz, Elke Sequence 19, CURRENT APPLICATION NUMBER: US/08/776,188B CURRENT FILING DATE: 2002-04-15 APPLICANT: Wiedmann, Michael TITLE OF INVENTION: DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING MULTIPLE ANTIGEN FILE REFERENCE: 100564-07003 Application US/08776188B Seidel, Christoph Ofenloch-Hahnle, Beatus

PRIOR APPLICATION NUMBER: P44 30 972.4

PRIOR APPLICATION NUMBER: P44 30 972.4

PRIOR APPLICATION NUMBER: P 44 30 972.4

PRIOR APPLICATION NUMBER: P 44 26 276.0

PRIOR APPLICATION NUMBER: P 44 26 276.0

PRIOR APPLICATION NUMBER: P 44 26 276.0

PRIOR FILING DATE: 1994-07-25

NUMBER OF SEQ ID NOS: 76

SOFTWARE: Patentin version 3.1

SEQ ID NO 19 FEATURE: OTHER INFORMATION: Description ORGANISM: Artificial Sequence of Artificial Sequence: SH-activated linear peptide

NAME/KEY: misc_feature LOCATION: (3)..(3) OTHER INFORMATION: XBB at position LOCATION: (2)..(2)
OTHER INFORMATION: Xaa at position 2 is Beta-alanine NAME/KEY: misc_feature w is Epsilon-aminocaproic acid

LOCATION: (4)...(4);
OTHER INFORMATION: XBB at US-08-776-188B-19 NAME/KEY: misc_feature position 4 is Beta-alanine

FEATURE:

12 LNSWGCKGRIICYTS 26 ß LSLWGCKGKLVCYTS 19

Best Local Similarity 66. Matches 10; Conservative

41.6%;

Score 72; Pred. No.

DB 4; 00066

Length 19

o;

Gaps

0;

Mismatches

Query Match Best Local 9

В

US-08-776-188B-36 Sequence 36, Application US/08776188B GENERAL INFORMATION:

APPLICANT:
APPLICANT: APPLICANT: Weinhues, Ursula-Henrike APPLICANT: Kruse-Muller, Cornelia APPLICANT: Hoss, Eva Faatz, Elke Ofenloch-Hahnle, Beatus

APPLICANT: Wiedmann, Michael TITLE OF INVENTION: DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING MULTIPLE ANTIGEN FILE REFERENCE: 100564-07003 CURRENT APPLICATION NUMBER: US/08/776,188B CURRENT FILING DATE: 2002-04-15 Seidel, Christoph

> NAME/KEY: misc_feature LOCATION: (4)..(4) OTHER INFORMATION: Xaa US-08-776-1888-36 SOFTWARE: PatentIn version 3.1 SEQ ID NO 36 LENGTH: 19 PRIOR FILING DATE: 1994-07-25
> NUMBER OF SEQ ID NOS: 76
> SOFTWARP: Date: 1994-07-25 PRIOR APPLICATION NUMBER: PCT/EP95/02919
> PRIOR FILING DATE: 1995-07-24
> PRIOR APPLICATION NUMBER: P 44 30 972.4
> PRIOR ETILING DATE: 1994-08-31
> PRIOR APPLICATION NUMBER: P 44 26 276.0
> PRIOR APPLICATION NUMBER: P 44 26 276.0 Query Match Best Local Similarity Matches LENGTH: 19 TYPE: PRT OTHER INFORMATION: Description OTHER INFORMATION: gp41/2 NAME/KEY: misc_feature LOCATION: (2)...(2)
> OTHER_INFORMATION: XAA at position LOCATION: (1)..(1)
> OTHER INFORMATION: Xaa at FEATURE: ORGANISM: Artificial Sequence NAME/KEY: misc_feature NAME/KEY: misc_feature LOCATION: (3)..(3) FEATURE: FEATURE: OTHER INFORMATION: Xaa at FEATURE: Conservative Xaa at position 4 is Beta-alanine 41.6%; position 1 is digoxigenin-3-cme position Score 72; Pred. No. of Mismatches Artificial Sequence: Digoxigenin-labelled linea ω N 18 ìs Epsilon-aminocaproic acid Beta-alanine 0.00066; DB 4; Length 19 Indels 0 Gaps

QΥ 밁 US-09-576-824A-20 12 LNSWGCKGRIICYTS 26 |: |||||:::|||| 5 LSLWGCKGKLVCYTS 19

0

CURRENT APPLICATION NUMBER: US/09/576,824A
CURRENT FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 08/723,425
PRIOR TILING DATE: 1996-09-30
PRIOR APPLICATION NUMBER: 09/146,028
PRIOR APPLICATION NUMBER: 09/146,028
PRIOR FILING DATE: 1993-11-29
PRIOR FILING DATE: 1993-03-08
PRIOR FILING DATE: 1993-03-08 Sequence 20, Applica GENERAL INFORMATION: APPLICANT: De Leys, R TITLE OF INVENTION: NUMBER OF SEQ ID NOS: 600 SOFTWARE: PatentIn Ver. 2. SEQ ID NO 20 LENGTH: 22 FILE REFERENCE: 2752-11 PRIOR APPLICATION NUMBER: EP 92400598.6 PRIOR FILING DATE: 1992-03-06 FEATURE: NAME/KEY: VARIANT ORGANISM: Human immunodeficiency virus Application US/09576824A PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR USE IN TO IMMUNOLOGICALLY IMPORTANT EPITOPES OF A PROCESS FOR DETERMINATION OF ANTIBODIES OF BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTED EPITOPES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM

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RESULT 11
US-08-776-188B-54
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                                                                                                                Sequence 54, Application US/08776188B
GENERAL INFORMATION:
APPLICANT: Weinhues, Ursula-Henrike
APPLICANT: Kruse-Muller, Cornelia
APPLICANT: Hoss, Eva
APPLICANT: Faatz, Elke
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Best Local ;
                                                                 APPLICANT:
                                                                                                       APPLICANT:
          CURRENT
                         APPLICANT: Wiedmann, Michael
TITLE OF INVENTION: DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING MULTIPLE ANTIGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Seidel, Christoph
APPLICANT: Wiedmann, Michael
TITLE OF INVENTION: DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING MULTIPLE ANTIGEN
FILE REFERENCE: 10564-07003
CURRENT APPLICATION NUMBER: US/08/776,188B
CURRENT APPLICATION NUMBER: US/08/776,188B
PRIOR APPLICATION NUMBER: P07-24
PRIOR FILING DATE: 1995-07-24
PRIOR PRIOR FILING DATE: 1995-07-24
PRIOR APPLICATION NUMBER: P 44 30 972.4
PRIOR APPLICATION NUMBER: P 44 30 972.4
PRIOR APPLICATION NUMBER: P 44 26 276.0
PRIOR FILING DATE: 1994-08-31
PRIOR APPLICATION NUMBER: P 44 26 276.0
PRIOR FILING DATE: 1994-07-25
PRIOR FILING DATE: 1994-07-25
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SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Description of Artificial Sequence: Epitope region of HIV I or OTHER INFORMATION: HIV subtype 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Weinhues, Ursula-Henrike APPLICANT: Kruse-Muller, Cornelia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
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OTHER INFORMATION: modified
NAME/KEY: VARIANT
LOCATION: (22)
                                                                                                                                                                                                                                                                                                                  9 QQRLNSWGCKGRIICYTSARW 29
| :||||| | :|:|:
| QAQLNSWGCAFRQVCHTTVPW 21
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APPLICATION NUMBER: US/08/776,188B
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                                                                                Ofenloch-Hahnle, Beatus
Seidel, Christoph
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Faatz, Elk
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Seidel, Christoph
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Pred. No. 0.00077;
2; Mismatches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                Length 27;
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RESULT 12
US-09-359-975-14
US-09-359-975-14
; Sequence 14, Application US/09359975
; GENERAL INFORMATION:
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LOCATION: (1)...(1); OTHER INFORMATION: Xaa at position 1 is BPRU
US-08-776-188B-54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 41.6%;
Best Local Similarity 52.4%;
Matches 11; Conservative
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LENGTH: 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 2002-04-15
PRIOR APPLICATION NUMBER: PCT/EP95/02919
PRIOR FILING DATE: 1995-07-24
PRIOR APPLICATION NUMBER: P 44 30 972.4
PRIOR FILING DATE: 1994-08-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Ruthenylated linear peptid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 QAQLNSWGCAFRQVCHTTVPW 25
                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 mb-MD/JAF
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/359,975
ETLING DATE: 23-Unl-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9 QQRLNSWGCKGRIICYTSARW 29
                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Weiner, David B. Williams, William V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                        APPLICATION NUMBER: 08/0 FILING DATE: 15-JUL-1993
                     APPLICATION NUMBER: 08/029,336 FILING DATE: 11-MAR-1993
                                                                                                    APPLICATION NUMBER: 08/124,962 FILING DATE: 21-SEP-1993
                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/979,385B
FILING DATE: 26-NOV-1997
APPLICATION NUMBER: US 08/495,684
FILING DATE: 28-SEP-1995
                                                                                                                                            APPLICATION NUMBER: 08/125,012 FILING DATE: 21-SEP-1993
                                                                                                                                                                                 APPLICATION NUMBER: PCT/US94/00899 FILING DATE: 26-JAN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Philadelphia
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & Norris STREET; One Liberty Place 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OF INVENTION: Compositions and Methods
Genetic Material
NUMBER: 08/008,342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bin
                                                                              08/093,235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 72;
Pred. No.
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RESULT 14
US-09-576-824A-6
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; OTHER INFORMATION: HIV subtype O
US-08-776-188b-5
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Sequence 6, Application US/09576824A GENERAL INFORMATION:
                                                                                                                                                                                            Matches
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Best Local
                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Wiedmann, Michael
TITLE OF INVENTION. DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING MULTIPLE ANTIGEN
FILE REFERENCE: 100564-07003
CURRENT APPLICATION NUMBER: US/08/776,188B
CURRENT FILING DATE: 2002-04-15
PRIOR APPLICATION NUMBER: PCT/EP95/02919
PRIOR APPLICATION NUMBER: PCT/EP95/02919
PRIOR APPLICATION NUMBER: P 44 30 972.4
PRIOR PILING DATE: 1994-08-31
PRIOR APPLICATION NUMBER: P 44 30 972.4
PRIOR PILING DATE: 1994-08-31
PRIOR APPLICATION NUMBER: P 44 30 972.4
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NUMBER OF SEQ ID NOS: 76
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 23
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Hoss, Eva APPLICANT: Faatz, Eli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Weinhues, Ursula-Henrike APPLICANT: Kruse-Muller, Cornelia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Deluca, Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: UPAP-0253
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEPA: 215-568-3429
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Watch 40.5%;
Local Similarity 52.4%;
                                                                                                                                                   12 LNSWGCKGRIICYTSARWH 30
                                                                                                                                                                                                            Local Similarity
                                                                                                               1 LGIWGCSGKLICTTAVPWN 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID
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Seidel, Christoph
                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30, Application US/08776188B GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: De Leys, R
TITLE OF INVENTION:
                                                                                                                                   SEQ ID NO 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                         PRIOR APPLICATION NUMBER: P 40 PRIOR FILING DATE: 1994-07-25 NUMBER OF SEQ ID NOS: 76
                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: PCT/EP95/02919
PRIOR FILING DATE: 1995-07-24
PRIOR APPLICATION NUMBER: P 44 30 972.4
PRIOR FILING DATE: 1994-08-31
                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/08/776,188B CURRENT FILING DATE: 2002-04-15
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Wiedmann, Michael TITLE OF INVENTION: DETERMINATION OF A SPECIFIC IMMUNOGLOBULIN USING MULTIPLE ANTIFILE REFERENCE: 100564-07003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 09/723,425
PRIOR FILING DATE: 1996-09-30
PRIOR APPLICATION NUMBER: 09/146,028
PRIOR APPLICATION NUMBER: 09/146,028
PRIOR FILING DATE: 1993-11-2
PRIOR APPLICATION NUMBER: PCT/EP93/00517
PRIOR FILING DATE: 1993-03-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                        LENGTH: 20
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 1992-03-06
NUMBER OF SEQ ID NOS: 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/576,824A CURRENT FILING DATE: 2000-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Weinhues, Ursula-Henrike
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: EP 92400598.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE
FEATURE:
OTHER INFORMATION: Description
OTHER INFORMATION: gp41/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: modified site NAME/KEY: VARIANT LOCATION: (24) OTHER INFORMATION: modified site
                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE:
                                                                                                                                                       PatentIn version
                                                                                                                                                                                                                                                                                                                                                                                                                                      Ofenloch-Hahnle, Beatus
Seidel, Christoph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hoss, Eva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kruse-Muller,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING PROCESS FOR THE IMPORTANT EPITOPES AND THEIR USE IN A PROCESS FOR DETERMINATION OF ANTIBODIES OF BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY EPITOPES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             32.1%;
47.8%;
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                        of Artificial Sequence:
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                        Digoxigenin-labelled linea
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HEATURE:

NAME/KEY: misc_feature

LOCATION: (1)...(1)

OTHER INFORMATION: Xaa at position 1 is digoxigenin-3-cme

FEATURE:

NAME/KEY: misc_feature

LOCATION: (2)...(2)

OTHER INFORMATION: Xaa at position 2 is Beta-alanine

FEATURE:

NAME/KEY: misc_feature

LOCATION: (3)...(3)

OTHER INFORMATION: Xaa at position 3 is Epsilon-aminocaproic acid

FEATURE:

NAME/KEY: misc_feature

LOCATION: (4)...(4)

COTHER INFORMATION: Xaa at position 4 is Beta-alanine

US-08-776-188B-30
Search completed: June 20, 2002, 15:34:57 Job time: 205 sec
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                                                                                                                                                     QУ
                                                                                                                                                                                               Query Match 31.8%; Score 55; DB 4; Length 20; Best Local Similarity 53.3%; Pred. No. 0.19; Matches 8; Conservative 3; Mismatches 4; Indels
                                                                                                             12 LNSWGCKGRIICYTS 26
| || |::|| |:
5 LGIWGCSGKLICTTA 19
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Page

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Title: Perfect score: Sequence:

US-09-605-573A-69 173

GRETLMQDQQRLNSWGCKGRIICYTSARWH

30

protein -

protein search, using sw model

GenCore version 4.5 Copyright (c) 1993 - 2000 Comp

Compugen

June 20,

2002,

Scoring |

table:

BLOSUM62

Gapop 10.0 ,

Gapext 0.5

Total number Searched:

of.

hits satisfying chosen parameters:

283138

283138 seqs, 96089334 residues

Maximum Minimum

DB seq

length: 0 length: 2000000000

Post-processing:

Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

PIR_71:*

pir1:*
pir2:*
pir3:*
pir4:*

Result No.

Score

Match Query

Length

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SUMMARIES

Pred. No. score grea and is der

668.91.00 668.91.00 667.91.00 667.91.00 667.

8 104 8 277 8 277 8 277 8 278

\$52930 \$49197 \$VCLJG3 VCLJG4 VCLJG4 VCLJG4 VCLJG7 VCLJG7 VCLJG17 VCLJS7 VCLJS7 VCLJS7 VCLJS7 VCLJS7 VCLJS7 VCLJS9 S30453 S30453 S30455 S30457 S30456 S30456 S30456 S30457 S30456 S30466 S304

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No. 1s the number of results predicted by chance to have a greater than or equal to the score of the result being printed, s derived by analysis of the total score distribution.
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(without alignments)
194.513 Million cell updates/sec
envelope protein g
envelope protein g
envelope protein g
envelope protein g
env polyprotein no
env protein - huma
env protein g
envelope protein g
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hrd
d
                                                                                                                                                 A;Status: preliminary
A;Molecule type: genomic RNA
A;Residues: 1-863 <VAN>
A;Cross-references: GB:L02587
C;Superfamily: type E retrovirus en
C;Keywords: polyprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1

S523930

C;Species: human immunodeficiency virus type 1 (fragment)

C;Species: human immunodeficiency virus type 1, HIV-1

C;Species: human immunodeficiency virus type 1, Change 26-Aug-1999

C;Date: 06-Jun-1995 #sequence_revision 21-Jul-1995 #text_change 26-Aug-1999

C;Chen, J.H.M.; Guetard, D.; Philbert, F.; Chamaret, S.; Tabary, T.; Montagnian submitted to the EMBL Data Library, January 1995

submitted to the EMBL Data Library Lib
                                                                                                                                                                                                                                                      A;Cross-references: EMBL:X84328; C;Superfamily: type E retrovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: genomic A; Residues: 1-104 <COH>
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A53034
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19; Consert
                                                                  ch 67.18;
1 Similarity 60.78;
17; Conservative
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67.9%;
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VCLJBR
T12016
S543916
S543984
S13288
VCLJA2
VCLJA2
VCLJA2
VCLJA3
VCLJAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NID:g695526; PIDN:CAA59066.1; PID:g695527
                                                                        ; Score 116; DB
; Pred. No. 1.7e
8; Mismatches
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pred. No. 1.2e-09;
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ETLLQNQQLLSLWGCKGKLVCYTSVKWN 613

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C; Species: human immunodeficiency virus type 1, HIV-1
C; Species: human immunodeficiency virus type 1, HIV-1
C; Date: 19-Mar-1997 **sequence_revision 24-Jul-1997 **text_change 26-Aug-1999
C; Accession: S49197
R; Charneau, p; Borman, A.M.; Quillent, C.; Guetard, D.; Chamaret, S.; Cohen, J.; Remy, submitted to the EMBL Data Library, July 1994
A; Description: Isolation and envelope sequence of a highly divergent HIV-1 isolate: defi A; Accession: S49197
A; Molecule type: DNA
A; Residues: 1-877 CCHAS
A; Cross-references: EMBL:X80020; NID:g510516; PIDN:CAA56323.1; PID:g510517
C; Superfamily: type E retrovirus env polyprotein
C; Keywords: glycoprotein; capsid protein; coat protein; fil-30/Domain: signal sequence **status predicted <CP1>F; 536-877/Product: coat protein gp120 **status predicted <CP1>F; 536-877/Product: coat protein gp41 **status predicted <CP1>F; 59,88,139,148,159,184,188,198,230,235,242,263,270,277,292,302,333,345,357,367,396,404,
                                                                                                                                                                                                                            C; Superfamily: type E retrovirus env polyprotein
C; Reywords: AIDS; capsid protein; coat protein; glycoprotein;
F: 130/Domain: signal sequence #status predicted <STG>
F: 31-500/Product: coat protein gp120 #status predicted <CP1>
F: 501-854/Product: coat protein gp41 #status predicted <CP2>
F: 501-854/Product: coat protein gp41 #status predicted <CP2>
F: 501-854/Product: transmembrane #status predicted <TM1>
F: 675-693/Domain: transmembrane #status predicted <TM2>
F: 675-693/Domain: transmembrane #status predicted <TM3>
F: 805-821/Domain: transmembrane #status predicted <TM3>
F: 134,140,143,154,158,186,195,239,260,267,274,299,331,336,351
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: EMBL:X52154; NID:g58866; PIDN:CAA36407.1; PID:g58874
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Species: simian immunodeficiency virus SIVcpz
A; Note: host Pan troglodytes (chimpanzee)
C; Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 16-Jul-1999
C; Accession: S09990
R; Huet, T; Cheynier, R; Meyerhans, A.; Roelants, G.; Wain-Hobson, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Gene: env
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: DNA
A; Residues: 1-854 <HUE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Status: nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Title: Genetic organization of a chimpanzee lentivirus related A;Reference number: S09983; MUID:90259077
A;Accession: S09990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N;Alternate names: coat polyprotein N;Contains: coat protein 9p120; coat protein 9p41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R:Huet, T.; Cheynier, R.; Meyerhans, A.; Roelants, Nature 345, 356-359, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           env polyprotein precursor - simian immunodeficiency
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S49197
envelope protein precursor - human immunodeficiency virus type 1 (fragment)
C;Species; human immunodeficiency virus type 1, HIV-1
C;Date: 19-Mar-1997 #sequence_revision 24-Jul-1997 #text_change 26-Aug-1999
C;Accession: S49197
A M.: Ouillent, C.; Guetard, D.; Chamaret, S.; Cohe
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                                      ETLMQDQQRLNSWGCKGRIICYTSARWH
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                                                                                                                                                Similarity
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18; Conservative
                                                                                                                        Conservative
                                                                                                                                           53.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            not shown; translation not shown
                                                                                                             Score 93; DB Pred. No. 3.2e 5; Mismatches
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Pred. No. 3
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3.2e-05;
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1.3e-08;
1es 5;
                                                                                                                                                                                                                       <TM3>
,331,336,351,356,384,392,426,432,446,45
                                                                                                               9,
                                                                                                                                                           Length 854
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                                                                                                                                                                                                                                                                                                                                                                                                        immunodeficiency; polyprot
                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to HIV-1.
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                                                                                                       Gaps
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envelope protein gp120/gp41 - human immunodeficiency virus type C;Species: human immunodeficiency virus type 1, HIV-1 A;Variety: isolate 20
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C;Superfamily: type E retrovirus env polyprotein
C;Keywords: AIDS; capsid protein; coat protein; glycoprotein; immunodeficiency; polyp
F;1-17/Domain: signal sequence #status predicted <SIG>
F;18-858/Product: env polyprotein #status predicted <MAT>
F;18-501/Product: exterior membrane glycoprotein #status predicted <EXT>
F;502-858/Product: transmembrane glycoprotein #status predicted <TMM>
F;504-858/Product: transmembrane glycoprotein #status predicted <FMM>
F;34,67,76,119,120,151,166,179,192,193,196,206,238,241,248,272,278,289,300,367,371,40
               В
                                                        Q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       env polyprotein precursor - human immunodeficiency virus type 2 (isolate ROD) N;Alternate names: coat polyprotein C;Species: human immunodeficiency virus type 2, HIV-2 C;Date: 30-Sep-1987 #sequence_revision 30-Sep-1987 #text_change 16-Jul-1999 C;Accession: C26262
                                                                                                                                                                                                                                                                                                                                                                                                   A;Gene: env
                                                                                                                                                                                                                                                                                                                                                                                                                    A; Cross-references: GB:M15390; NID:g1332361; PIDN:AAB00770.1; PID:g325749
                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-858 <GUY>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Contents: proviral DNA
A; Accession: C26262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Title: Genome organization and transactivation of the human A;Reference number: A26262; MUID:87173056
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Guyader, M.; Emerman, M.; Sonigo, Nature 326, 662-669, 1987
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                                                                                                         Matches
                                                                                                                                                  Query Match
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                                              3 ETLMQDQQRLNSWGCKGRIICYTSARW 29
EKYLQDQARLNSWGCAFRQVCHTTVPW 609
                                                                                                      L Similarity
15; Conserv
                                                                                                   Conservative
                                                                                                                        53.2%;
                                                                                                                     Score 92;
Pred. No.
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                                                                                              Mismatches
                                                                                  DB 1;
. 4.4e-05;
8;
                                                                                                                                       Length 858
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                                                                                         Indels
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                                                                                  Gaps
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RESULT
S21990
                                                                                                                                                                                                                                    A;Gene: env
C;Superfamily: type E retrovirus (
C;Keywords: AIDS; capsid protein;
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A;Title: Sequence of simian immunodeficiency virus from African green monkey, a new m A;Reference number: A30045; MUID:88232906
A;Accession: G30045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C:Species: simian immunodeficiency viru
C:Date: 30-Jun-1989 #sequence_revision
C:Accession: G30045
                                                                                                                                                                                                                                                                                                                        A; Cross-references: EMBL: X07805
                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
A; Residues: 1-864 <FUK>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              env polyprotein - simian immunodeficiency virus (African green monkey isolate)
c;Species: simian immunodeficiency virus, SIV
c;Date: 30-Jun-1989 #sequence_revision 30-Jun-1989 #text_change 24-Oct-1997
                                                                                                                                                                        Query Match
Best Local
                                                                                                                                                      Matches
                                                                          809
                                                                                                      3 ETLMQDQQRLNSWGCKGRIICYTSARW 29
                                                                 EKYLEDQARLNSWGCAWKQVCHTTVEW
                                                                                                                                                                  Similarity
                                                                                                                                                  Conservative
                                                                                                                                                                51.4%;
48.1%;
                                                                                                                                                                                                                                        env polyprotein
; coat protein;
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Pred. No.
                                                                                                                                           Mismatches
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J.00012;
8;
                                                                                                                                                                                                                                    immunodeficiency;
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envelope protein gp120/gp41 · human immunodeficiency virus type 1 c:Species: human immunodeficiency virus type 1, HIV-1 A;Variety: isolate 3L C:Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 01-1 C:Accession: S22002; S70418 C:Accession: S22002; S70418 R:Steuler, H.; Storch-Hagenlocher, B.; Wildemann, B.; Hacke, W. submitted to the EMBL Data Library, July 1991 A;Description: Distinct populations of HIV-1 in blood and cerebrosp A;Reference number: S21990 A;Accession: S22002 A;Molecule type: DNA A;Residues: 1-358 <STEIL SA;Cross-references: EMBL:X61352; NID:g60186; PIDN:CAA43616.1; PID:g R;Steuler, H.; Storch-Hagenlocher, B.; Wildemann, B.
                                                                                              RESULT
S22000
                                                                                                                                                                                                                                                                                                                                                                               A; Reference number: $70417; MUID:92144209
A; Accession: $70418
A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-333, 'X', 335-358 <STE2>
A; Cross references: EMBL:X61352; NID:g60186
C; Superfamily: type E retrovirus env polyprotein
envelope protein gp120/gp41 - human immunodeficiency virus type 1
C:Species: human immunodeficiency virus type 1, HIV-1
C:Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 12-Apr-1995
C:Accession: S22000
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C;Accession: S21990; S70423
R;Steuler, H; Storch-Hagenlocher, B.; Wildemann, B.;
submitted to the EMBL Data Library, July 1991
A;Description: Distinct populations of HIV-1 in blood
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AIDS Res. Hum. Retroviruses 8, 53-59, 1992
A:Tille: Distinct populations of human immunodeficiency virus type 1 in blood A:Reference number: S70417; MUID:92144209
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A;Residues: 1-332,'X',334-357 <STE2>
A;Cross-references: EMBL:X61357; NID:g60175; PIDN:CAA43626.1; PID:g60176
C;Superfamily: type E retrovirus env polyprotein
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AIDS Res. Hum. Retroviruses 8, 53-59, 1992
A;Title: Distinct populations of human immunodeficiency A;Reference number: S70417; MUID:92144209
A;Accession: S70423
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A; Residues: 1-357 <STE1>
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Best Local S
Matches 14
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Matches 14
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$21990
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Pred. No. 7.2e
5; Mismatches
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Pred. No.
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mann, B.
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                                                                                                                                                                                                                                                                       .2e-05;
9;
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R;Steuler, H; Storch
submitted to the EMBI
A;Description: Distin
A;Reference number: &
A;Accession: $22000
C:Superfamily: type E retrovirus env polyprotein C:Keywords: coat protein; glycoprotein; polyprotein; tran: F:1-19/Domain: signal sequence #status predicted <SIG>F:20-859/Product: env polyprotein #status predicted <ENV>F:20-501/Product: surface glycoprotein gp120 #status predicted <ENV>
                                                                                                                                                                                   A;Title: Molecular characterization of an A;Reference number: A33943; MUID:90112662 A;Accession: H33943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: genomic RNA
A; Residues: 1-859 <KUM>
                                                                                                                                                                                                                                       C;Date: 31-Dec-1990 #sequence_revision 31-Dec-1990 #text_change 09-Sep-1994 C;Accession: H33943 R;Kumar, P.; Hui, H.; Kappes, J.C.; Haggarty, B.S.; Hoxie, J.A.; Arya, S.K.; J. Virol. 64, 890-901, 1990
                                                                                                                                                                                                                                                                                                                              N;Contains: surface glycoprotein gp120;
C;Species: human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A: Molecule type: DNA
A: Residues: 1-358 <STE>
A: Cross-references: EMBI
C: Superfamily: type E ro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       envelope protein gp120/gp41 - human immunodeficiency virus type C;Species: human immunodeficiency virus type 1, HIV-1 A;Variety: patient 3B C;Date: 19-Mar-1997 *sequence_revision 25-Apr-1997 *text_change C;Accession: S70417
                                                                                                                 C; Genetics:
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                                                                                              A; Gene:
                                                                                                                                  A; Cross-references:
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C;Superfamily: type
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A; Residues: 1-358 <STE>
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Best Local (
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Best Local Similarity
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type E retroviru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Storch-Hagenlocher, B.; Wildemann, he EMBL Data Library, July 1991
Distinct populations of HIV-1 in bluber: S21990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           type E retrovirus
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Pred. No. 0.00
4; Mismatches
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0.0001;
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   predicted
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A;Cross references: EMBL:M87071
A;Experimental source: FOENVA3
R;Gao, F.; Yue, L.; White, A.T.; Pappas, P.G.; Barchue, J.; Hanson, A.P.; Greene, B.M.; submitted to the EMBL Data Library, December 1992
A;Description: Human infection by genetically diverse SIVSM-related HIV-2 in west Africa A;Reference number: S30460
A;Accession: S30480
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C;Date: 02-Dec-1993 #sequence_revision 30-7an-1998 #text_change 17-Mar-1999
C;Accession: S30448; S30480; S30481
R;Gao, F.; Yue, L.; White, A.T.; Pappas, P.G.; Barchue, J.; Hanson, A.P.; G.
Nature 358, 495-499, 1992
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                   env protein - human immunodeficiency virus type 2 (fragment) C;Species: human immunodeficiency virus type 2, HIV-2 C;Date: 02-Dec-1993 #sequence_revision 30-Jan-1998 #text_cha C;Accession: S30453
R;Gao, F; Yue, L; White, A.T.; Pappas, P.G.; Barchue, J.;
                                                                                                                                            RESULT
S30453
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C;Superfamily: type E retrovirus env polyprotein
C;Keywords: coat protein; glycoprotein
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A; Residues: 1-151 <GAW>
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A;Accession: S30448
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A; Experimental source: FOENVAl3
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A; Residues: 1-151 <GA3>
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A;Accession: S30481
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R;Gao, F.; Yue, L.;
Nature 358, 495-499,
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Pred. No. 6.
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                                                                       #text_change 17-Mar-1999
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A; Status: translation not shown A; Molecule type: nucleic acid A; Residues: 1-151 <GAO>
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C;Species: human immunodeficiency virus type 2, HIV-2
C;Date: 02-Dec-1993 #sequence_revision 30-Jan-1998 #text_cha
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C;Keywords: coat protein; glycoprotein
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A;Title: Human infection by genetically diverse SIV(SM)-related HIV-2 in West Afri A;Reference number: S30448; MUID:92350299
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C;Accession: S30452
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A;Gene: env
C;Superfamily: type E retrovirus env polyprotein
C;Keywords: coat protein; glycoprotein
                                                                                                           A;Status: translation not shown
A;Molecule type: nucleic acid
A;Residues: 1-151 <GAO>
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A;Tille: Human infection by genetically diverse SIV(SM).
A;Reference number: S30448; MUID:92350299
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48.1%;
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Pred. No. 6.2e-05;
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Pred. No. 6.
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	20, 2002, 15;32:12	atch 49.7%; Score 86; DB 2; Le cal Similarity 48.1%; Pred. No. 6.2e-05; 13; Conservative 6; Mismatches 8; ETLMQDQQRLNSWGCKGRIICYTSARW 29 :: : : : : : EKYLKDQAQLNSWGCAFRQVCHTTVEW 54
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Listing first 45 summaries
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        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1 GRETLMQDQQRLNSW
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ENV_HV1S1
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P12449 human
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P05880 human
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P19559 human
P19549 human
P19551 human
P19551 human
P03378 human
P03378 human
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P04586 human
P04578 human
P05877 human
P05877 human
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Q74126 human immun
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AC P17281;
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N EINNED (GECNOC:) (FOIENITAE).	.:	N-LINKED (GLCNAC) (POTENTIAL).	(GLCNAC) ((GLCNAC)	(GLCNAC)	(GLCNAC:)	(GLCNAC)	(GLCNAC)	N-LINKED (GEOMAC:) (FOIENTEDE).	(GLCNAC)	CICNAC .	POTENTIAL.	POTENTIAL.	TRANSMEMBRANE GLYCOPROTEIN.	EXTERIOR MEMBRANE GLYCOPROTEIN.		Polyprotein; Glycoprotein; Transmembrane;			-	P41.				sb-sib.ch).	agreement (See http://w	is not removed. Usage by and for commerc	itions as long as its content is in no	Bioinformatics Institute. There are no restrictions on its	produced through		ENV POLYPROTEIN.	'	chimpanzee lentivirus related to E	A Roelants G Wain-Hobson S	100136			Retroviridae: Lentivirus.		smembrane glycoprotein (GP41)].	precursor (Contains: F		sequence update)	(t)));

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ENV_HYZKR
ID ENV_H
AC 07412
DT 01-NO
DT 01-NO
DT 01-NO
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                                                                                                                                                                                                                                                                                                                                                                                      use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kraus G.K., Talbott R., Leavitt M., Luznick L., Schmidt A., Badel P., Bartz C., Morton W., Wong-Staal F., Looney D.J.; Submitted (APR-1995) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Envelope polyprotein GP160 precursor [Contains: Exterior membrane
                                                                                                                                                                                                                                                                             Pfam; PF00516; GP120; Pfam; PF00517; GP41; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Viruses; Retroid viruses; NCBI_TaxID=73484;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus type 2 (isolate KR) (HIV-2). Viruses; Retroid viruses; Retroviridae; Lentivirus.
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                 (POTENTIAL).
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ID ENV_HVZRO
AC P04577;
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DT 13-AUG-1987
DT 16-OCT-2001
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                                 AIDS; Co
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13-AUG-1987 (Rel. 05, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Envelope polyprotein GP160 precursor [Contains: Exterior membrane glycoprotein (GP120); Transmembrane glycoprotein (GP41)].
ENV.
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                                                                                                                                                                                    use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                    MEDLINE-87173056; PubMed-3031510; Fine Median M., Sonigo P.,
                                                                                                                                  EMBL; M15390; AAB00770.1;
EMBL; X05291; CAA28914.1;
PIR; C26262; VCLJG2.
HIV; M15390; ENV$2ROD.
                                                                                                                                                                                                                                                         This
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                                                                                                  Pfam;
                                                                                                                                                                                                                        the European Bioinformatics Institute use by non-profit institutions as 10
                                                                                                                                                                                                                                                                               Nature 326:662-669(1987)
                                                                                                                                                                                                                                                                                                                                                                    Viruses; Retroid
NCBI_TaxID=11720;
                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus type 2 (isolate ROD) (HIV-2).
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InterPro; IPR000777; GP120.
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immunodeficiency virus type
                                                                                                                                                                                                                                                                                                                 Alizon M.;
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TRANSMEMBRANE GLYCOPROTEIN.
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N-LINKED (GLCNAC. . . ) (POI
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                                              GLYCOPROTEIN
                                                                              Transmembrane;
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01-NOV-1988 (Rel. 09, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Envelope polyprotein GP160 precursor [Contains:
glycoprotein (GP120); Transmembrane glycoprotein
                                          the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and fo entities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                SEQUENCE FROM N.A.
MEDLINE-8823906; Pubmed-3374586;
Fukasawa M., Miura T., Hasegawa A.,
Miki K., Kitamura T., Hayami M.;
                                                                                                                                                                                                                                                                                                                   Simian immunodeficiency virus (TYO-1 isolate) (SIV-AGM)
                                                                                                                                                                                                                                                                                     Viruses; Retroid
NCBI_TaxID=11731;
                                                                                                                                                        "Sequence of simian immunodeficiency virus from African gre
new member of the HIV/SIV group.";
Nature 33:457-461(1988).
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G30045; VCLJG4.
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EKYLODQARLNSWGCAFRQVCHTTVPW 609
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sen the Swiss Institute of Bioinformatics and the EMBL
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15; Conser
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Pred. No. 3.9e
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CRC64;
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Matches 13
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01-JUL-1993 (Rel. 26, Last seque
16-OCT-2001 (Rel. 40, Last annot:
Envelope polyprotein GP160 precui
glycoprotein (GP120); Transmembra
                                                                                                                                            SEQUENCE FROM N.A.

MEDLINE-91220680; PubMed-2024476;
Fomsgaard A., Hirsch V.M., Allan J.

"A highly divergent proviral DNA cl

of African green monkey.";

Virology 182:397-402(1991).

VIROLOGY 182:397-402(1991).
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              the European Bioinformatics Institute. use by non-profit institutions as lomodified and this statement is not remo
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InterPro; IPR000328; Env_GP41.
InterPro; IPR000777; GP120.
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                                                                             SWISS-PROT entry is copyright. It is produced through a collaboration sen the Swiss Institute of Bioinformatics and the EMBL outstation
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SP160 precursor [Contains: E
Transmembrane glycoprotein
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RESULT 6
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Watches 13; Conserv
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SEQUENCE FROM N.A.

MEDLINE-88320359; PubMed=3261862;

MEDLINE-88320359; PubMed=3261862;

Zagury J.F., Franchini G., Reitz M.S. Jr., Collalti E., Starcıcıı
Hall L., Fargnoli K., Jagodzinski L.L., Guo H.-G., Laure F.,
Hall L., Fargnoli K., Jagodzinski L.L., Gollo R.C.;

Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

"Arya S.K., Josephs S.F., Zagury D., Wong-Staal F., Gallo R.C.;

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01-NOV-1988 (Rel. 09, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Envelope polyprotein GP160 precursor [Contains: Exterior membrane glycoprotein (GP120); Transmembrane glycoprotein (GP41)].
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NCBI_TaxID=11719;
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Pfam; PF00516; Gp120; 1.
Pfam; PF00517; GP41; 1.
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viruses; Retroviridae; Lentivirus.
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P12449;
P1-0CT-1989 (Rel. 12, Created)
O1-OCT-1989 (Rel. 12, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
Envelope polyprotein GP160 precursor [
glycoprotein (GP120); Transmembrane gl
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Viruses; Retroid
NCBI_TaxID=11718
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InterPro; IPR000777; GP120.
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J03654; ENV$2NIHZ
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14; Conservative
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5; Mismatches
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RESULT 8
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SEQUENCE FROM N.A.
MEDLINE-89184641; PubMed-2648404;
Franchini G., Fargnoli K.A., Giombini F.,
Bosch M., Biberfeld G., Fenyo A.M., Albert
Bosch M.; Biberfeld G., Fenyo A.M., Albert
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Signal
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HIV; J04498; ENV$2ISY.
InterPro; IPR000328; Env_GP41.
InterPro; IPR000777; GP120.
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Pfam; PF00517;
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Proc. Natl. Acad. Sci. U.S.A. 86:2433-2437(1989).
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t J., Gallo R.C.
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MEDLINE-89228766; PubMed-2713163;
MEDLINE-89228766; PubMed-2713163;
Srinivasan A., York D., Butler D. Jr., Jannoun-Nasr McCormick J., Ou C.Y., Myers G., Smith T., Chen E.;
"Molecular characterization of HIV-1 isolated from a "Molecular characterization of minus of the molecular characterization of minus of the minus of the molecular characterization of minus of the minus of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a c between the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and fo entities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; M15896; AAB53948
PIR; A44963; A44963.
HIV; M15896; ENV$Z321.
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generation of hybrid HIV.";
AIDS Res. Hum. Retroviruses 5:121-129(1989).
                                                                                                                                                                                                                                      InterPro; IPR000328; Env_GP41.
InterPro; IPR000777; GP120.
Pfam; PF00516; GP120; 1.
Pfam; PF00517; GP41; 1.
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Viruses; Retroid viruses; Ret
NCBI_TaxID=11692;
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                                                                                                                                                                      Coat protein; Polyprotein; Glycoprotein; Transmembrane;
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Retroviridae; Lentivirus.
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GLYCOPROTEIN
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01-JUN-1994 (Rel. 29, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Envelope polyprotein GP160 precursor [Contains: Exterior membrane glycoprotein (GP120); Transmembrane glycoprotein (GP41)].
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Shaw
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                                                                                                                                                                                                                                                                                                                                                                                                        "Complete nucleotide sequence, genome organization, properties of human immunodeficiency virus type 1 i for limited defectiveness and complementation.";
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NCBI_TaxID=36377;
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InterPro; IPR000777; GP120.
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Retroviridae; Lentivirus
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MEDLINE-90347835; PubMed=2384920;
MEDLINE-90347835; PubMed=2384920;
Cheng-Mayer C., Quiroga M., Tung J.W., Dina D., Levy
Cheng-Mayer C., Quiroga M., Tung J.W., Dina D., Levy
"Viral determinants of human immunodeficiency virus
"Viral determinants of human immunodeficiency virus
"Viral determinants of human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                      01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Envelope polyprotein GP160 precursor [Contains: Exterior membrane glycoprotein (GP41)].
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InterPro; IPR000777; GP120.
                                                                                                                                                                                                                        macrophage tropism, cytopathogenicity, J. Virol. 64:4390-4398(1990).
                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                               NCBI_TaxID=11691;
                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus type 1 (SF162 isolate) (HIV-1).
Viruses; Retroid viruses; Retroviridae; Lentivirus.
                                                                                                                EMBL; M65024; AAA45072.1;
HIV; M38428; ENV$SF162.
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13; Conserv
                                                                                                                                             an email to license@isb-sib.ch).
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46.4%;
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   EXTERIOR MEMBRATE GTRANSMEMBRANE GBY SIMILARITY.
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                                  MEMBRANE
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4.4e-05;
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DT _01-NO

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DE _Bryvel

DE _91ycc

GN _ENV.

OS _Human

OC _V1rue

OX _NCBI_
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RX _MEDL1

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MEDLINE-86235450; PubMed-3012778;

MEDLINE-86235450; PubMed-3012778;

MEDLINE-86235450; PubMed-3012778;

MEDLINE-86235450; PubMed-3012778;

MEDLINE-86235450; PubMed-3012778;

Gallo R.C., Par

"Genetic variation in HTLV-III/LAV over time in at risk for AIDS.";

science 232:1548-1553(1986).

"I- MISCELLANEOUS: ISOLATES WMJ1, WMJ2, AND WMJ BLOOD SAMPLES SEQUENTIALLY TAKEN FROM A TWO WAS PERINATALLY INFECTED BY HER MOTHER.
                                                     This SWI
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENV_HYIW2 STANDARD; PRT; 847 AA. P05880; 01-NOV-1988 (Rel. 09, Created) 01-NOV-1988 (Rel. 09, Last sequence update) 15-JUL-1999 (Rel. 38, Last annotation update) Envelope polyprotein GP160 precursor [Contains: Eglycoprotein (GP120); Transmembrane glycoprotein
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11705;
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                   SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation - European Bioinformatics Institute. There are no restrictions on its
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4.4e-05;
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RESULT 12

ENV_HV1B8 STANDARD; PR:

ID ENV_HV1B8 STANDARD; PR:

AC P04582;

DT 13-AUG-1987 (Rel. 05, Created)

DT 13-AUG-1987 (Rel. 05, Last sequent of the 
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         Transmembrane glycoprotein (GP41)]
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MEDLINE-8511123; PubMed=2578615;

MATHET L., Haseltine W., Patarca R., Livak K.J., Starcich I

Josephs S.F., Doran E.R., Rafalski J.A., Whitehorn E.A.,

Baumelster K., Ivanoff L., Petteway S.R. Jr., Pearson M.L.,

Lautenberger J.A., Papas T.S., Ghrayeb J., Chang N.T., Gali
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HIV; K02011; ENV$BH8.
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01-OCT-1989 (Rel. 12, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Envelope polyprotein GP160 precursor [Contains: Exterior membrane glycoprotein (GP120); Transmembrane glycoprotein (GP41)].
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HIV; M21098; ENV$BRVA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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es; Retroid viruses; Retroviridae; Lentivirus.
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Best Loc
Matches
                                         EMBL; M38427; AAA45067.1; -.
HIV; M38427; ENV$SF33.
InterPro; IPR000328; Env_GP41.
InterPro; IPR000777; GP120.
Pfam; PF00516; GP120; 1.
Pfam; PF00517; GP41; 1.
AIDS; Coat protein; Polyprotein.
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O1-FEB-1991 (Rel. 17, Created)

O1-FEB-1991 (Rel. 17, Last sequence update)

16-OCT-2001 (Rel. 40, Last annotation update)

Envelope polyprotein (GP160) precursor (Contains: E:
91ycoprotein (GP120); Transmembrane glycoprotein
                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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SEQUENCE FROM N.A.

MEDLINE=90317906; PubMed=2370688;

York-Higgins D., Cheng-Mayer C., Bauer D., Levy J.A., Din:
York-Higgins D., Cheng-Mayer S., Bauer D., Levy J.A., Din:
"Human immunodeficiency virus type 1 cellular host range,
"human immunodeficiency virus type 1 cellular host range,
"human immunodeficiency virus type 1 cellular host range,
       Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                 replication, and cytopathicity the viral genome.";
J. Virol. 64:4016-4020(1990).
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NCBI_TaxID=11690;
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
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                                       Polyprotein; Glycoprotein;
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 Stevenson M., Haggerty S., Lamonica C., Mann A.M., Meier C., Wasiak A., "Cloning and characterization of human immunodeficiency virus variants diminished in the ability to induce syncytium-indepency tolysis.";
                                                                                                               Human immunodeficiency virus type 1 (MFA isolate) (HIV-1).
Viruses; Retroid viruses; Retroviridae; Lentivirus.
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           syncytium-independent
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           ETLMQDQQRLNSWGCKGRIICYTSARWH 30
 ERYLKDQQLLGIWGCSGKLICTTAVPWN 609
                                   ch 49.1%;
1 Similarity 46.4%;
13; Conservative
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Search completed: June

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ALIGNMENTS

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SULT 2	Qy 3 ETLMQDQQRLNSWGCKGRIICYTSARWH	Query Match Best Local Similarity 78. Matches 22; Conservative	SQ SEQUENCE 161 AA; 19	Transmembrane.	Pfam; PF00517; GP41;		RT "Phylogenetic analysis RL Submitted (JAN-1999) t	RC STRAIN-BCF57;	RP SEQUENCE FROM N.A.	OC Viruses; Retroid viruses; Retrovi		DT 01-DEC-2001 (TremBirel.	01-OCT-2000		SULT 1
	GRIICYTSARWH 30 - : :	76.9%; Score 133; DB 15; Length 161; 78.6%; Pred. No. 7.5e-13; ive 4; Mismatches 2; Indels 0; Gaps 0;	61 19305 MW; SE3AF197E1FDEE7C CRC64;	1	1.	253.1;	"Phylogenetic analysis and sub-typing of 47 HIV-1 group O isolates."; Submitted (JAN-1999) to the EMBL/GenBank/DDBJ databases.	 D., Sandrine S., Christel D., Francois S.,		Viruses; Retroid viruses; Retroviridae; Lentivirus.		1. 19, Last sequence update)	15, Creat	RY; PRT; 161 AA.	

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AIDS;
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InterPro; IPR000777; GP120.
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Roques P., Robertson
Philippe M.;
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Viruses; Retroid viruses; Retroviridae; Lentivirus
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Pfam; PF00517; GP41;
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EMBL/GenBank/DDBJ data
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Pred. No. 1.2e
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Roques P., Rol
Philippe M.;
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                                                                                                Roques P., Robertson D., Damond F., Sousquiere Depienne C., Brun-Vezinet F., Dormont D.;
"HIV-1 group O phylogenetic analysis of C2-gp4 Submitted (FEB-1999) to the EMBL/GenBank/DDBJ EMBL; AJ133056; CAB96217.1; ...
InterPro; IPR000727; GP120.
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pfam; PF00517; GP41; 1.
ALDS; Coat protein; Glycoprotein; Polyprotein;
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Viruses; Retroid viruses; Retroviridae;
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78.6%;
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Last
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                                                                                                                                                                                                                                                                                                                                                                                                          CRC64;
                                                CRC64
                                                                                                                                                                                                                                                                                                                                                                             Length 195;
                    Length 576;
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                  Query Match
Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 75.7
Best Local Similarity 75.0
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN-GROUP O:
STRAIN-GROUP O:
Bibollet-Ruche F., Ekasa E., Peeters M.,
"Molecular characterization of envelope t
14 new human immunodeficiency virus type
41fferent African countries";
                                                                                                                                                                                        NON_TER
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                            different African countries.";
different African countries.";
Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; Y09772; CAA70911.1;
InterPro; IPR00032B; Env_GP41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             O40456

O40456;
O1-JAN-1998 (TrEMBLrel. 05, Created)
O1-JAN-1998 (TrEMBLrel. 05, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ENVELOPE TRANSMEMBRANE GLYCOPROTEIN (FRAGMENT).
Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
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                                                                                                                                                                                                                                                                                          Transmembrane.
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Q9IEC8;
Q9IEC8;
01-OCT-2000 (TrEMBLrel. 15,
01-OCT-2000 (TrEMBLrel. 15,
01-DEC-2001 (TrEMBLrel. 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Phylogenetic analysis and sub-typing of 47 HIV-1 group O isolates."; Submitted (JAN-1999) to the EMBL/GenBank/DDBJ databases. EMBL; AJ36391; CAB96240.1; -. InterPro; IPR000328; Env_GP41. Pfam; PF00517; GP41; 1. Transmembrance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fransmembrane.
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Philippe M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN-BCF02;
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nilarity 75.08;
Conservative
                                                                                                                                                                                     114
114 AA;
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219
219 AA;
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                                                                                                                                                                             114
; 13737 MW;
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219
; 25363 MW;
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75.0%;
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   Score 130; DB 15;
Pred. No. 1.5e-12;
4; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Created)
Last sequence update)
Last annotation update)
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Pred. No. 2.
                                                                                                                                                                          7907F77F5B51FA9D CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     85C2DC5B5F528907 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               s M., Delaporte E.;
lope transmembrane glycoprotein
type 1 group O strains from
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?.1e-12;
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                                                                    Length 114;
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       Indels
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   0,
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Q9IEC5
    Query Match 74.6%;
Best Local Similarity 71.4%;
Matches 20; Conservative
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Best Local Similarity
Matches 20; Conserv
                                                                                                                           NON_TER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q9IEB6;
01-OCT-2000 (TrEMBLrel. 15,
01-OCT-2000 (TrEMBLrel. 15,
01-DEC-2001 (TrEMBLrel. 19,
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NON_TER 1

NON_TER 216

SEQUENCE 216
                                                                                                        SEQUENCE
                                                                                                                                                                 Transmembrane.
                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae;
NCBI_TaxID=11676;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             GP41 (FRAGMENT).
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                                                                                                                                                                                                                                                                                       Philippe M.
                                                                                                                                                                                                                                                                                                          Roques P.,
                                                                                                                                                                                                                                                                                                                             STRAIN-BCF14;
                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Viruses; Retroid
NCBI_TaxID=11676;
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01-DEC-2001
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219 i
                                                                                                                                                                                                                                                                                                     Robertson
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216
216 AA;
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viruses; Retroviridae;
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25353
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216
; 25027 MW;
                                                                                                                                                                                                                                                                                                   D., Sandrine S.,
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19,
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Score 129; DB 15;
Pred. No. 4.4e-12;
6; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Created)
Last sequence update)
Last annotation update)
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Pred. No. 4.36
6; Mismatches
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Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                  F65829EB01520D91 CRC64;
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                                                                                                                                                                                                                                                                                           Christel D., Francois
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 15,
4.3e-12;
2;
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                                                                                                                                                                                                                                                           47 HIV-1
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databases.
                                     Length
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Best Local S
Matches 21
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O11940;
O11940;
O11940;
O1-JUL-1997 (TrEMBLrel. 04, Created)
O1-JUL-1997 (TrEMBLrel. 04, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ENVELOPE GLYCOPROTEIN (FRAGMENT).
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SEQUENCE
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Q9IHU1;
01-OCT-2000 (
01-OCT-2000 (
01-DEC-2001 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE-97340911; PubMed-9197385;
Brennan C.A., Hackett J. Jr., Zekeng
Hickman R.K., Gurtler L., Kaptue L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Devare S.G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-ABT1123;
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ELYCLOPE GLYCOPROTEIN (FRAGMENT).
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pfam; PF00517; GP41; 1.
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EMBL; U90133; AAB62816.1; -.
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                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus type 1. Viruses; Retroid viruses; Retroviridae;
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                                                                                                                                                                         Yang C., Gao F., Fonjungo P.N., Zekeng
Pieniazek D., Schable C., Lal R.B.;
"Phylogenetic analysis of protease and
                                                                            type 1 group O.";
AIDS Res. Hum. Retroviruses 16:1075-1081(2000)
EMBL; AF229243; AAF71920.1; -.
InterPro; IPR000328; Env_GP41.
                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. STRAIN=97CM747;
  NON_TER
                                                                                                                                                                                                                                        MEDLINE=20386754; PubMed=10933623;
                                              Transmembrane.
                                                                   Pfam; PF00517; GP41;
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75.0%;
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Pred. No.
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Overbeck J., Hampl
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Best Local Similarity 75.0
Matches 21; Conservative
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Best Local
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Q9WRV5;
01-NOV-1999
01-NOV-1999
01-DEC-2001
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NON_TER
SEQUENCE
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Transmembrane.
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Mas A., Quinones-Mateu M.E., Domingo E., Soriano
"Phylogeny of HIV type 1 group 0 isolates based o
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                                          AIDS Res.
EMBL; AF0
                                                                         MEDLINE-99284364; PubMed=10357472;
MAS A., Quinones-Mateu M.E., Domingo E., S
"Phylogeny of HIV type 1 group O isolates
sequences.";
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Viruses; Retroid viruses; Retroviridae;
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01-DEC-2001
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NCBI_TaxID=11676;
                     InterPro;
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ETLIQNQORLDLWGCKGRIICYTSVKWN 59
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Res. Hum. Retroviruses 15:769-773(1999).
AF081815; AAD41227.1; -.
Pro; IPR000328; Env_GP41.
PF00517; GP41; 1.
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(TrEMBLrel. 12,
(TrEMBLrel. 19,
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(TrEMBLrel. 12,
(TrEMBLrel. 19,
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75.0%;
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Last annotation updat
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Last sequence update)
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Pred. No. 3
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No. 3.7e-12;
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RESULT 14
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DT 01-NOV
OS HUMBAN
OC VIRUSE
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Best Local Similarity 71.4%;
Matches 20; Conservative
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SEQUENCE FROM N.A.

STRAIN-ABT063;

MEDLINE-97340911; PubMed-9197385;

Brennan C.A., Hackett J. Jr., Zekeng L., Lund J.K., Vallari A.S.,
Hickman R.K., Gurtler L., Kaptue L., von Overbeck J., Hampl H.,
Devare S.G.;

"Sequence of gp4lenv immunodominant region of HIV type 1 group 0 from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Sim
Matches 21;
                                                                                                                                                                                                                                                                                                                     Ol1939 PRELIMINARY; PRT; 105 AA.
011931:
01-JUL-1997 (TrEMBLrel. 04, Created)
01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ENVELOPE GLYCOPROTEIN (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   J. Virol. 68:1581-1585(1994).
EMBL; L2057; AAA448641.;
InterPro; IPR000328; Env_GP41.
InterPro; IPR000777; GP120.
Pfam; PF00516; GP120; 1.
Pfam; PF00517; GP41; 1.
AIDS; Coat protein; Glycoprotein
AIDS; Coat protein; Glycoprotein
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079670; OT-NOV-1996 (Tremburel 01, Cr
01-NOV-1996 (Tremburel 01, La
01-DEC-2001 (Tremburel 19, La
ENVELOPE PROTEIN GP120/GP41.
                                                                                                                                                                                                                           Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus
NCBI_TaxID=11676;
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SEQUENCE FROM N.A.

MEDLINE=94149848; PubMed=8107219;

Gurtler L.G., Hauser P.H., Eberle J., von Brunn A., Knapp S.,

Zekeng L., Tasque J.M., Kaptue L.;

A new subtype of human immunodeficiency virus type 1 (MVP-5180)
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Viruses; Retroid viruses; Retroviridae; Lentivirus
NCBI_TaxID=11676;
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SEQUENCE 143 AA; J
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||||:|:|||||||||:|||||:||
|96 ETLIONOORLNLWGCKGKLICYTSVKWN 623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 ETLMODOQRLNSWGCKGRIICYTSARWH 30
|||:|:|:| || ||||||:|||| ||:
61 ETLIONOQLLNLWGCKGRIVCYTSVRWN 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 75.0
21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein; Glycoprotein; Polyprotein; Transmembrane.
876 AA; 99245 MW; A92F868E37522BAB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16331 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      74.0%;
75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Score 128; DB; Pred. No. 2.8e
6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Created)
Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ٠.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 128; DB 15;
Pred. No. 3.9e-12;
4; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15A17063CCD3431D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 15;
2.8e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ş
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 876;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 143;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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В
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                                                                                                RT DR DR DR DR FT FT SQ
                                        Query Match 73.4%;
Best Local Similarity 71.4%;
Matches 20; Conservative
                                                                                             Transmembrane.
NON_TER 105
NON_TER 105
SEQUENCE 105 A
                                                                                                                        west central Africa.";
AIDS Res. Hum. Retroviruses 13:901-904(1997).
EMBL; U90132; AAB62815.1; -.
InterPro; IPR000328; Env_GP41.
Pfam; PF00517; GP41; 1.
105
105 AA;
                                                                                        1
105
; 12531 MW;
                                       Score 127; DB 15;
Pred. No. 4e-12;
5; Mismatches 3;
                                                                                    2D3D6B5BADC4A382 CRC64;
                                       ω
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                                                       Length 105;
                                    Indels
                                    0,
                                  Gaps
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Search completed: June 20, 2002, 15:34:36 Job time: 214 sec